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## ALIGNMENTS

RESULT 1
ADM16603
ID ADM1 WPI; 2004-270022/25. Cytostatic; nootropic; neuroprotective; antiparkinsonian; inhibitor; cell migration; cell invasion; cell proliferation; angiogenesis; apoptosis; tumour growth rate; human kininogen domain 3v; HK-D3v; HK neurodegenerative disease; Alzheimer's disease; Parkinson's disease; Donate 13-SEP-2002; 2002US-0410279P 15-SEP-2003; 2003WO-US028881. 25-MAR-2004. WO2004024877-A2 Synthetic. Homo sapiens. amyotrophic lateral sclerosis; ALS; mutein. HK-D3v(GS) 17-JUN-2004 ADM16603; ADM16603 standard; protein; 125 (ATTE-) ATTENUON LLC **'**3 Mazar amino acid sequence, (first entry) ð seq ₽ id Ŋ..

HK-D3;

Novel isolated anti-angiogenic polypeptide comprising sequence of human kininogen domain 3v (HK-D3v) or variant of native HK-D3, useful for inhibiting angiogenesis, endothelial cell proliferation or endothelial tube formation.

N-PSDB; ADM16606.

Disclosure; SEQ ID NO 2; 79pp; English.

The invention relates to an isolated anti-angiogenic polypeptide (I) comprising the sequence of human kininogen domain 3v (HK-D3v) (fully defined sequence (S1) of 127 amino acids as given in specification) or variant or derivative of native HK-D3 (fully defined sequence (S2) of 123 amino acids as given in specification), and has at least 20% of activity of native HK-D3 in inhibiting angiogenesis, endothelial cell inhibiting angiogenesis, endothelial cell

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC diagnostically or therapeutically labelled anti-angiogenic polypeptide (II) comprising (I) labelled with a diagnostic or therapeutic label, and CC (II) comprising (I) labelled with a diagnostic or therapeutic label, and CC with a detectable label and a carrier. Further disclosed is an affinity CC ligand (IX) useful for binding to or isolating an HK-D3-binding molecule CC or cells expressing the binding molecule, comprising (II) immobilized to a correct or carrier. (I) or (II) are useful for inhibiting cell conducing apoptosis which involves contacting cells associated with CC inducing apoptosis which involves contacting cells associated with CC undesired cell migration, invasion, proliferation or angiogenesis with CC (I) is useful for treating neurodegenerative diseases e.g., (I) is useful for treating neurodegenerative diseases e.g., (IX) is useful for treating neurodegenerative diseases e.g., (LS). The current sequence represents an N-terminal addition variant of terminal of the N-terminal addition variant of terminal collegenerative contacting an additional GS at the N-terminal collegenerative contacting collegenerative contacting cells as sociated with collegenerative contacting cells as collegenerative contacting cells as collegenerative diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                       protein engineering; kininogen; antiangiogenic; apoptosis; angiogenesis; cancer; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Variant human kininogen domain 3 Hk-D3v(GS) protein.
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                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                              Novel human kininogen domain 3 variant inhibiting angiogenesis, endothelial cell proliferation or endothelial tube formation, treating disease associated with undesired cell migration, invo
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                                                                                                 N-PSDB;
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proliferation or angiogenesis

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cc variant or derivative, a variant or derivative of native HK-D3 having a cfully defined 123 amino acid sequence given in the specification, which completely compared to the activity of polypeptide, variant or derivative has at least 20% of the activity of native HK-D3 in inhibiting angiogenesis, endothelial cell proliferation compared to a magiogenesis, endothelial cell proliferation compared to a magiogenesis, or for inducing apoptosis, which involves contacting comparising (I) is useful for therapeutically labeled (I). A composition comparising (I) is useful for treating subject having a disease or comparising associated with undesired cell migration, invasion, proliferation comparising associated with undesired cell migration, invasion, composition composition composition composition associated with undesired cell migration, invasion, composition compos
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Anti-angiogenic; angiogenesis; inhibitor; kininogen; homologue; endothelial cell proliferation; apoptosis; cancer; ocular disor rheumatoid arthritis; cytostatic; antiarthritic; antirheumatic;
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                      WPI; 2000-442247/38
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                                                                                                                                                                                                                                                                                                                                                                                                                           human; D3 peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               r; ocular disorder; antirheumatic; therapy;
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Novel isolated anti-angiogenic polypeptide comprising sequence of human kiningen domain 3v (HK-D3v) or variant of native HK-D3, useful for
                                                                                                          N-PSDB; ADM16605.
                                                                                                                                                                                                                                                                                                                    13-SEP-2002; 2002US-0410279P
                                                                                                                                                                                                                                                                                                                                                                            15-SEP-2003; 2003WO-US028881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytostatic; nootropic; neuroprotective, antiparkinsonian; inhibitor; cell migration; cell invasion; cell proliferation; angiogenesis; apoptosis; tumour growth rate; human kininogen domain 3v; HK-D3v; HK neurodegenerative disease; Alzheimer's disease; Parkinson's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                           (ATTE-) ATTENUON LLC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amyotrophic lateral sclerosis; ALS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Native HK-D3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Composition for inhibiting angiogenesis and endothelial cell proliferation, inducing endothelial cell apoptosis and treating cancer, rheumatoid arthritis, and ocular disorders comprises a kininogen domain 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 GKDFVQPPTKICVGCPRDIPTNSPELBETLTHTITKLNAENNATFYFKIDNVKXARVQVV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYFTVNCQF
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                                                                                                                                                                                                  Mazar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98.5%;
100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 123;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HK-D3v; HK-D3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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CC variant or derivative of native HK-D3 (fully defined sequence (82) of 123 cc amino acids as given in specification), and has at least 20% of activity cc of native HK-D3 in inhibiting angiogenesis, endothelial cell ccl proliferation or endothelial tube formation. Also disclosed is a cc diagnostically or therapeutically labelled anti-angiogenic polypeptide (II) comprising (I) labelled with a diagnostic or therapeutic label, and a diagnostic HK-D3-related composition (III) comprising (II) labelled with a diagnostic or therapeutic label, and detectable label and a carrier. Further disclosed is an affinity cl jand (IX) useful for binding to or isolating an HK-D3-binding molecule cor cells expressing the binding molecule, comprising (I) immobilized to a solid support or carrier. (I) or (II) are useful for inhibiting cell ingration, cell invasion, proliferation or angiogenesis, or for undesired cell invasion, invasion, proliferation or angiogenesis with (I) (I) is useful for treating neurodegenerative diseases e.g., Alzheimer's disease, Parkinson's disease or amyotrophic lateral sclerosis cequence.
Sequence 123 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to an isolated anti-angiogenic comprising the sequence of human kininogen domain 3v defined sequence (S1) of 127 amino acids as given in variant or derivative of native HK-D3 (fully defined
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tube formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inhibiting angiogenesis, endothelial cell proliferation or endothelial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              79pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                        specification) or specification of 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polypeptide
(HK-D3v) (fi
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Best Local :
121
                  123
                                                                                                                   123;
                                      61
                                                         63
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                                                                                                                              Similarity
                   203
                                 AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
                                                                                    GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                 AGKKYFIDFVARETTCSKESNEELTESCETKKLGOSLDCNAEVYVVPWEKKIYPTVNCQP
                                                                        GKDFVQPPTK1CVGCPRD1PTNSPELEETLTHT1TKLNAENNATFYFK1DNVKKARVQVV
                                                                                                                    Conservative
123
                  125
                                                                                                              98.5%; but
100.0%; Pr
                                                                                                                         Score 660; DB 8; Pred. No. 4.4e-66;
                                                                                                                   Mismatches
                                                                                                                                    Length 123;
                                                                                                                Indels
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                                                                                                                Gaps
                                   120
                                                      122
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                                                                                             62
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RESULT
ADVI195
ID ADD
XX AD
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Donate F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein engineering; cancer; cytostatic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2005058599-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Native human kininogen domain 3 protein
                                                                                                    (DONA/) DONATE F. (MAZA/) MAZAR A P.
                                                                                                                                                                                                                                                                 15-SEP-2003; 2003US-00661784
                                                                                                                                                                                                                                                                                                                                                                15-SEP-2003; 2003US-00661784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADY71956;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADY71956 standard; protein; 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-JUN-2005
Mazar AP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    kininogen; antiangiogenic; apoptosis; angiogenesis;
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WPI;

2005-213544/22

for

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The invention relates to an isolated anti-antiangiogenic polypeptide (I) the sequence of human kininogen domain 3 variant (HK-D3v) having a cc fully defined 127 amino acid sequence given in the specification, or its call and the specification of the comparing a cc variant or derivative, a variant or derivative of native HK-D3 having a cc fully defined 123 amino acid sequence given in the specification, which cc polypeptide, variant or derivative has at least 20% of the activity of cc native HK-D3 in inhibiting angiogenesis, endothelial cell proliferation cc or endothelial tube formation in an in vitro or in vivo bioassay. (I) is correspond for inhibiting cell migration, cell invasion, cell proliferation cc angiogenesis, or for inducing apoptosis, which involves contacting cc angiogenesis with (I) or a therapeutically labeled (I). A composition cc cangiogenesis with (I) or a therapeutically labeled (I). A composition condition associated with undesired cell migration, invasion, proliferation cc condition associated with undesired cell migration, invasion, contacting condition associated with undesired cell migration, invasion, contacting condition associated with undesired cell migration, invasion, contacting cc condition associated with undesired cell migration, invasion, invasion, cc condition associated with undesired cell migration, invasion, invasion, cc condition associated with undesired cell migration, invasion, invasion, cc condition associated with undesired cell migration, invasion, invasion, cc condition associated with undesired cell migration. Accomposition cc condition associated with undesired cell migration, invasion, providing to a cc condition associated with undesired cell migration. Accomposition cc condition associated with undesired cell migration, invasion, contacting cc condition associated with undesired cell migration. This sequence cc condition associated with undesired cell migration and contacting condition and contacting condition and contacting condition and condition and conditio
RESULT 6
ADD43683
ID ADD84
XX ADP4
AC ADP4
XX 18-1
XX 18-1
XX 19-1
XX bum
KW 93-8
KW 93-8
KW 93-8
KW 10-1
XX 10-1
XX 10-1
XX 10-1
XX WO:
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XX PN WO:
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Best Local S
Matches 123
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                                                                                                                                                                                                                                                                 gastrointestinal; cardiovascular; immunosuppressive; antiinflammatory; cytostatic; neuroprotective; gynaecological; gene therapy; cytostatic; neuroprotective; cardiovascular disorder; autoimmune disorder; gastrointestinal disorder; cardiovascular disorder; autoimmune disorder; inflammatory disorder; cell proliferative disorder; disorder; developmental disorder; epithelial disorder; neurological disorder; developmental disorder; epithelial disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human pMMM-40 protein SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADP43683 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                                         human; protein modification and maintenance molecule; PMMM;
                                                                                                                            WO2004053068-A2
                                                                                                                                                                                                                                                 reproductive
                                                                    24-JUN-2004
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ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                        disorder.
               2003WO-US038573.
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; Pred. No. 4.4e-66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NO:40
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05-DEC-2002;
17-DEC-2002;
24-JAN-2003;
30-JAN-2003;
21-FEB-2003;
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Kable AE,
                                                                                                                                                                                                                                             (INCY-) INCYTE CORP.
                                                                                                                                                                                                                  WPI; 2004-468843/44.
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o, Blake
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2003US-0444141P.
2003US-0449491P.
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2002US-0434315P.
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Khare R, Lee SY, Ti
Ho A, Zheng W, Gao
                                                                                                                                                                                                                                         Ramkumar
                                                                                                                                                                                                                                  Tran UK,
Gao J, Che
                                                                                                                                                                                                                                  J, Richardson TW;
UK, Bhatia UG;
Chawla NK, Mason
                                                                                                                                                                                                                                    Mason PM;
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New human protein modification and maintenance molecules and polynucleotides for diagnosing, preventing or treating diseases polynucleotides for diagnosing preventing or creating diseases associated with aberrant protein expression, e.g. cardiovascular or cell associated with aberrant protein expression, e.g. proliferative disorders.

**莱星**莱星星星素器莱克克克克克克克克克克克克克克克克克克克克克克克

Claim 1; SEQ ID NO 40; 268pp; English.

maintenance molecule (PMMM) (I), designated pmMM-40. Also described: (1) cmaintenance molecule (PMMM) (I), designated pmodification and comprising a promoter sequence operably linked to the polynucleotide in a isolated polynucleotide in comprising a plutransformed with the recombinant polynucleotide in comprising a plutransformed with the recombinant polynucleotide (1); (2) a cellar polynucleotide in comprising a plutransformed with the recombinant polynucleotide (2); (3) a cellar polynucleotide in comprising the precombinant polynucleotide or (1); (3) a cellar polynucleotide or (1); (3) a cellar polynucleotide or (1); (3) a cellar polynucleotide or (1); (4) a composition or an attained proposition or compound, an antagonist compound or an antibody, and an exciptent; (5) or treating diseases or conditions associated with decreased expression or treating diseases or condition or an antibody, and an exciptent; (5) that is effective as an agonist or antigonist of (1), or is effective in that is effective as an agonist or antigonist of (1), or is effective in that is effective as an agonist or antigonist of (1), or is effective in condition or disease associated with the expression of the target polynucleotide; (11) perpenting a polyclonal or or disease associated with the expression or monoclonal antibody with a subject; (14) preparing a polyclonal or monoclonal antibody with the tribody produced by the method in (14); (15) a composition are monoclonal antibody and a subject; (14) preparing a polyclonal polyclonal or monoclonal antibody with the subject of a sample containing the compositions or polyclonal antibody. And a subject is a composition profile of a sample containing the compositions are polyclonal antibody. And a subject is a contained to a contained the effect of the polymelotide molecule composition and methods are useful for compositions and can be effective. The composition a protein-protein interactions,

268 AA

Query Match Best Local Similarity Conservative 98.5%; Score 660; DB 8; 100.0%; Pred. No. 1.3e-65; Mismatches Length 268 Indels 0

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RESULT 7
ADM83721
ADM932
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The invention relates to novel diagnostic and therapeutic polynucleotides

Selected from one of the 2722 sequences defined in the specification. A

polynucleotide of the invention may have a use in gene therapy. The human

diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be

used to diagnose a particular condition, disease or disorder associated

with human molecules, e.g. cell proliferative disorders, or

disorder, neurological disorders, gastrointestinal disorders, or

disorder, neurological disorders, gastrointestinal disorders, or

infections caused by virus, bacteria, fungi or parasite. The dithp

molecules may also be used in genetic mapping, in identifying individuals

from minute biological samples, in detecting single nucleotide

polymorphisms, as molecular weight markers, and for somatic or germline

gene therapy. The present sequence represents a dithp protein of the

cuther therapy. The sequence data for this patent is not represented in

the printed specification, but was obtained in electronic format directly
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lagace RE, Span-
Xu Y, Kwong M, Po
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mooney EM, I
Stevens KA,
Peralta CH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in diagnosing a condition, molecules, e.g. autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schmidt JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-SEP-2002; 2002US-0410259P 12-SEP-2002; 2002US-0410260P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human diagnostic and therapeutic pprotein SEQ ID NO:3970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Harthshorne TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-SEP-2003; 2003WO-US028227
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diagnostic and therapeutic polynuclectides and polypeptides, useful diagnosing a condition, disease or disorder associated with human scules, e.g. autoimmune or inflammatory disorders, in gene therapy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2004-329368/30.
DB; ACN42373.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page; 190pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin
Anderson SB, Rioux P, Shen EJ, Wick, Stuve LL;
Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton
ng M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, G
Shi X, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wright RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   suchorolski MT,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnostic and therapeutic polynucleotide; dithp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pitte SJ, bic. Pitte SJ, Bic. TP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Elder LV;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kirton ES;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-AUG-2001;
14-DEC-2001;
18-JAN-2002;
                                                                                                                                                                                     New human extracellular messenger (EXMES) polypeptide, useful for preparing a composition for treating a disease associated with decreased expression or overexpression of functional EXMES e.g. autoimmune
                                                                                                                                                                                                                                                                                                                                                                                  Ding L, yao m
Richardson TW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-MAR-2002;
05-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; extracellular messenger; EXMES; cytostatic; antidiabetic; immunosuppressive; autoimmune disorder; inflammatory disorder; diabetes; endocrine disorder; cancer.
                                                                                                                                                                                                                                                                                                                                                              Ramkumar J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABP70801
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                                                                                                                                                                                                                                                                                                                                                                                                                                   SY,
                                                                                                                                                                                                                                                                                        2003-278643/27.
DB; ACC42388.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INCYTE GENOMICS
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                                                                                                                                                                                                                                                                                                                                                                                                        Yao MG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lee S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001US-0314811P.
2001US-0340584P.
2002US-0350595P.
2002US-0363432P.
2002US-03634607P.
2002US-0370761P.
2002US-0391378P.
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                                                                                                                                                                                                                                                                                                                                                                               R, Wilson AD, Jin P,
G, Becha SD, Tran UK,
Kable AE, Chang H, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein; 304
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Baughn MR,
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Pred. No. 1.3e-65;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hafalia AJA,
                                                                                                                                                                                                                                                                                                                                                                               , Hawkins PR, Tang YT, B
, Chien D, Zebarjadian Y;
Swarnakar A, Lehr-Mason P
                                                                                                                                                                                                                                                                                                                                                                                                                                           Walia NK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 268;
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The present invention relates to novel human extracellular messenger proteins (EXMES-1 to-28; ABP70774-ABP70801) and their coding sequence (ACC42361-ACC42388). The proteins are useful for preparing a composit for diagnosing or treating a disease or condition associated with

sequences composition

Page

224pp;

English

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RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                          Lagace RE,
Xu Y, Kwon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stevens KA,
Peralta CH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dith) or polypeptides may be used to diagnose a particular condition, disease or disorder associated with human molecules, e.g. cell proliferative disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mooney EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Schmidt JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-SEP-2002;
12-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-SEP-2003; 2003WO-US028227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2004023973-A2
                                                                                                                                                                                                                                             in diagnosing a condition, molecules, e.g. autoimmune
                                                                                                                                                                                                                                               New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Harthshorne TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (INCY-) INCYTE CORP.
                                                                                                                                                                           Claim 27;
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DB; ACN42372.
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                                                                                                                                                                                                                                                                                                                                                                                                            t JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F;
shorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV;
r EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP;
r EKA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH;
r EKA, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
r RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES
Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietz
r S, Shi X, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                         mapping
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                                                                                                                                                                           Page; 190pp;
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[larity 100.0%;
Conservative
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2002US-0410260P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and therapeutic pprotein SEQ ID NO:3969
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                                                                                                                                                                           English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stuve LL;
UA, Kirton ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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RESULT 10
ABP70799
ID ABP70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  autoimmune/inflammatory disorder, developmental disorder, endocrine disorder, neurological disorders, gastrointestinal disorders, or infections caused by virus, bacteria, fungi or parasite. The dithp molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline gene therapy. The present sequence represents a dithp protein of the invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                                        24-AUG-2001; 2001US-0314811P.
14-DEC-2001; 2001US-0340584P.
18-JAN-2002; 2002US-0350595P.
11-MAR-2002; 2002US-0363432P.
15-MAR-2002; 2002US-0370761P.
05-APR-2002; 2002US-0391378P.
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                                                                                                                                                                                                                                                                                                                                        immunosuppressive; autoimmune
endocrine disorder; cancer.
                                                                                                                                                                                                                                                                                                                                                    Human; extracellular messenger; EXMES; cytostatic; antidiabetic; immunosuppressive; autoimmune disorder; inflammatory disorder; c
                                                                                                                                                                                                                                                                                                                                                                                      Human extracellular messenger,
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                                                                                                   Duggan
Lee SY,
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                                                                                                                                                                                                                                                                                              WO2003018612-A2
                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                             Ding L, Yao M
Richardson TW,
                                                                                                                                                                                                                                                 22-AUG-2002; 2002WO-US027213
                                 N-PSDB;
                                                                                                                                     (INCY-)
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                               2003-278643/27.
)B; ACC42386.
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                                                                                                                                     INCYTE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
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                                                                                      Khare R,
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ilarity 100.0%;
Conservative
                                                                                              Lee S,
hare R,
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                                                                                                                                       GENOMICS
                                                                              Kable AE
                                                                            , Baughn MR, Hafal
Wilson AD, Jin P,
Becha SD, Tran UK,
able AE, Chang H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein;
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                                                                                                                                       INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            322
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Pred. No. 1.5e-65;
                                                                                , Hafalia AJA, Wa
Jin P, Hawkins P
Tran UK, Chien D,
ang H, Swarnakar A
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                                                                                           walia NK, Elliot
B PR, Tang YT, B
D, Zebarjadian Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 304
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                                                                                                                    Elliott VS;
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                                                                                    짬.
                                                                                                        Burford
                                                                                                                                                                                                                                                                                                                                                           diabetes;
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New human extracellular preparing a composition

messenger (EXMES) polypeptide, useful for treating a disease associated with

with

decreased

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RESULT 11
ABM83717
ID ABM83717
ID ABM83717
ID ABM83717
ID ABM837X
AC ABM83

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Best Local S
Matches 123
                                                                                                                                                                                                                                                                  Harthshorne :
Mooney EM, I
Stevens KA,
Peralta CH,
New diagnostic and therapeutic polynuclectides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy of in gene mapping.
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Xu Y,
                                                                                                                                                                                                                                                                                                                                                                               Schmidt JP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human diagnostic and therapeutic pprotein SEQ ID NO:3966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-NOV-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to novel human extracellular messenger proteins (EXMES-1 to-28; ABB700774-ABB70801) and their coding sequences (ACC42361-ACC42388). The proteins are useful for preparing a compositic for diagnosing or treating a disease or condition associated with decreased expression or overexpression of functional EXMES e.g. autoimmune/inflammatory disorders, diabetes, endocrine disorders or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disorders
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                                                                                                                                                2004-329368/30.
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                                                                                                                                                                                            RE, Spiro PA, Stewar
Kwong M, Policky JL,
S, Shi X, Suarez CJ;
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                                                                                                                                                                                  Wright RJ, Bruns CM, Marjanovic ...,
TA, Suchorolski MT, Altus CM, Pitts SJ, Elder Lv;
Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin,
Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Splro PA, Stewart EA, Wingrove J, Vitt UA, Kirtor
Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirtor
Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirtor
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2002US-0410260P
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                                                                                                                                                                                                                                                                                                                                                                                                                                CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnostic and therapeutic polynucleotide; dithp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98.5%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
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                                                                                                                                                                                                                                                                                            Gerstin EH;
                                                                                                                                                                                                                                                Kirton ES;
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Best Local Similarity
                       12-OCT-2001;
25-OCT-2001;
09-NOV-2001;
28-NOV-2001;
06-DEC-2001;
14-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell proliferation disorder; atherosclerosis; neurological disorder; epileps; Huntington's disease; stroke; immune disorder; allergy; inflammatory disorder; AIDS; developmental disorder; hypothyroidism; Cushing's syndrome; gastrointestinal disorder; epithelial disorder; infection; cytostatic; antiarteriosclerotic; anticonvulsant; nootropinfection; cytostatic; nootropinfection; nootropinfection; nootropinfection; nootropinfection; nootropi
                                                                                                                                                                                                                       11-OCT-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disorder, neurological disorders, gastrointestinal disorders, or infections caused by virus, bacteria, fungi or parasite. The dithp molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline gene therapy. The present sequence represents a dithp protein of the invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; protein modification and maintenance molecule;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human protein modification and maintenance molecule-24 (PMMM-24).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-JUL-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU92044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABU92044 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     used to diagnose a particular condition, disease or diso with human molecules, e.g. cell proliferative disorders, autoimmune/inflammatory disorder, developmental disorder autoimmune/inflammatory disorder, developmental disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      326 AA;
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                    ; 2001US-0329689P.
; 2001US-0335703P.
; 2001US-03348887P.
; 2001US-0334145P.
; 2001US-0337451P.
; 2001US-0337451P.
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                                                                                                                                                                                                                          2002WO-US032850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cerebroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  thyromimetic.
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271 122 211

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The present invention relates to the isolation of human protein CC modification and maintenance molecules (PMMM), and the polynucleotide CC sequences encoding them. A total of 40 pMMM polypeptides (designated pMMM CC = 1 to pMMM-40) are disclosed. The sequences of the invention are useful CC for diagnosing a condition or disease associated with the expression of CC pMMM in a subject, preparing a polyclonal or monoclonal antibody, and CC generating an expression profile of a sample containing the CC expression or overexpression of pMMM are cell proliferation disorders (e.g. epilepsy, CC (e.g. cancer, atherosclerosis), neurological disorders (e.g. epilepsy, CC (e.g. cancer, atherosclerosis), neurological disorders, (e.g. epilepsy, CC allergies), developmental disorders (e.g. hypothyroidism, Cushing's CC allergies), developmental disorders (e.g. hypothyroidism, Cushing's pMMM polypeptides or their fragments are useful in screening compounds (e.g. epilepsy), correspondents are useful in screening compounds that the activity of the polypeptides, or in CC altering the expression of the target polynucleotide and compounds that general polynucleotide and compounds that perform the polypeptides or that the activity of the polypeptide.

ABU92021-ABU92060 represent the human pMMM polypeptides of the invention
                                                                                                                                                                                                                                RESULT 13
ABM83716
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Best Local S
Matches 123
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Tran UK, Becha SD, Duggan BM, Lee EA, Griffin JA, Li JX;
Sprague WW, Hafalia AJA, Chawla NK, Lehr-Mason PM, Kable AE, Marquie JP, Yao MG, Richardson TW, Tang TY, Jin P, Chien D;
Bhatia U, Burrill JD, Lee S, Blake JJ, Ho A, Zheng W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New human protein modification and maintenance molecules (PMMM), useful for diagnosing, treating and preventing diseases or conditions associativith the aberrant PMMM expression e.g. cancer, atherosclerosis, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-430274/40.
N-PSDB; ACA92439.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 264-265; 311pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 329 AA;
                                                                                          gene therapy;
                                                                                                                                                                                                                     ABM83716 standard; protein;
                                                                                                                            Human
                                                                                                                                                           18-NOV-2004
                               WO2004023973-A2
                                                                                                                                                                                                                                                                                                                                                                                                                             155
                                                                                                                                                                                                                                                                                                                                 123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          123;
                                                                                                                       diagnostic and therapeutic pprotein SEQ ID NO:3965
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                                                                                                                                                                                                                                                                                                                                   LGM 125
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Pred. No.
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                                                                                                polynucleotide; dithp
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ABP70800
ID ABP7
XX
AC ABP7
XX
DT 26-J
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26-AUG-2003 ABP70800; ABP70800

(first

entry)

Human extracellular messenger, EXMES-27

RESULT 14

standard; protein;

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The invention relates to novel diagnostic and therapeutic polynucleotides composited from one of the 2722 sequences defined in the specification. A complex condition of the invention may have a use in gene therapy. The human composition and therapeutic polynucleotides (dithp) or polyneptides may be used to diagnose a particular condition, disease or disorder associated composition with human molecules, e.g. cell proliferative disorder, endocrine autoimmune/inflammatory disorder, developmental disorder, or composition disorder, or composition caused by virus, bacteria, fungion parasite. The dithp composition caused by virus, bacteria, fungion parasite. The dithp composition in the bological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline composition. Note: The sequence represents a dithp protesin of the composition. Note: The sequence represents a dithp protesin of the composition of the printed specification, but was obtained in electronic format directly from wife at www.wipo.int/pct/en/sequences/listing.htm
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-SEP-2002; 2002US-0410259P.
12-SEP-2002; 2002US-0410260P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-SEP-2003; 2003WO-US028227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 27; Page; 190pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-329368/30.
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                                                                                                                                                                                                                                                            Sequence 329
                                                                                                                                                                                                            Local
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                             123
                                                             215
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S, Shi X,
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                                                          AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
                                                                                                                             GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ght RJ, Bruns CM, Marjanovi
Suchorolski MT, Altus CM,
                                                                                                                                                                                                                 Score 660;
Pred. No.
                                                                                                                                                                                                   Mismatches
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CM, Pitts SJ, Blder L
CM, SC, Reddy TP,
                                                                                                                                                                                                                   1.7e-65;
                                                                                                                                                                                                                                      DB 8;
                                                                                                                                                                                                                                  Length 329;
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Human protein #41

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18-JAN-2002; 2002US-0350595P.
11-MAR-2002; 2002US-0363432P.
15-MAR-2002; 2002US-0364607P.
05-APR-2002; 2002US-0370761P.
24-JUN-2002; 2002US-0391378P.
              08-SEP-2005
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                                          AEB32339,
                                                                    AEB32339 standard;
                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                     for diagnosing or treating a disease or condition associated wi decreased expression or overexpression of functional EXMES e.g. autoimmune/inflammatory disorders, diabetes, endocrine disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 206; 224pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disorders or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New human extracellular messenger (EXMES) polypeptide, useful fo preparing a composition for treating a disease associated with despression or overexpression of functional EXMES e.g. autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB;
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Richardson
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                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to novel human extracellular messenger proteins (EXMES-1 to-28; ABP70774-ABP70801) and their coding sequences (ACC42361-ACC42388). The proteins are useful for preparing a composition for diagnosing or treating a disease or condition associated with
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SY, Khare R, Wilson AD, Jin P, Hawkins PR, Tang YT, Bux
g L, Yao MG, Becha SD, Tran UK, Chien D, Zebarjadian Y,
hardson TW, Kable AE, Chang H, Swarnakar A, Lehr-Mason PM,
                                                                                                                                                                  123
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                                                                                                                                                                                          AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
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             (first entry)
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                                                                  protein;
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                                                                                                                                                                                                                                                                                                                 98.5%; Score 660; DB 6; 1
100.0%; Pred. No. 1.9e-65;
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AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 122 AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP

307

GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV

247

248

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188

Query Match Best Local S Matches 123

Similarity

98.5%;

Score 660; DB 9; Pred. No. 1.9e-65;

Length 362; Indels

123;

Conservative

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Mismatches

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Gaps

Sequence

362

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The invention relates to a detection reagent capable of detecting one or composed mucleic acid polymorphisms. The invention also relates to compose determining whether a trait is linked to one of the human chromosomes or compately readable medium having stored in it the SNP concession, a computer readable medium having stored in it the SNP concession at least about 12 contiguous mucleotides, genotyping at least concession in a sample, identifying an comprising at least about 12 contiguous mucleotides, genotyping at least concession in a sample, identifying an comprising at least one container containing the detection reagent. CC the sub-region comprises determining whether a trait is linked to one of the human chromosomes or its sub-region comprises determining whether that differentiates between alternative composition given in the specification in a sample comprises contacting the detection reagents. Genotyping at least one SNP position given in the specification given in the specification. CC alleles at at least one SNP position given in the specification. CC alleles at at least one SNP position given in the specification in a concess of determining which allele is present at the at least one SNP position. CC comprises genotyping at least one SNP specification in a nucleic acid sample from the individual. The disorder is type II diabetes. CC comprises genotyping at least one SNP specification in a cc concession and diabetes or obesity. This sequence concession in a sequence data for this patent did not form part of the printed cc sequence data for this patent did not form part of the printed cc sequence data for this patent did not form part of the printed cc sequence attained in the scope of the invention. Note: The sequence attained in the scope of the invention. Supposed the sequence in the scope of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New detection reagent capable of detecting 1, 100, 500, 1000 or 5000 or more single nucleic acid polymorphisms, useful in identifying an individual having or at risk of developing type II diabetes or obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 102; 31pp; English.
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10-SEP-2001; 2001US-00948947.
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DB; AEB32278.
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Maximum DB
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length: 2000000000
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Match
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1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/6_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/FCTUS_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
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Copyright (c) 1993 - 2006 Biocceleration Ltd
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US-08-791-522-1
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US-09-461-061-4
US-09-461-061-4
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Sequence 480, App
Sequence 29, Appl
Sequence 475, App
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Sequence 1, Appli
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## ALIGNMENTS

US-09-461-061-18 RESULT 1

US/09461061

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TITLE OF INVENTION: Inhibition of Angiogenesis By TITLE OF INVENTION: Kininogen Domain 3 Peptide An. FILE REFERENCE: 6056-260 US
CURRENT APPLICATION NUMBER: US/09/461,061
CURRENT FILING DATE: 1999-12-15
EARLIER APPLICATION NUMBER: 60/112,427
EARLIER FILING DATE: 1998-12-16
NUMBER: 0F SEQ ID NOS: 21
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 18
LENGTH: 123
RESULT 2
US-09-949-002-476
(S-09-949-002-476)
; Sequence 476, Application US/09949002
; Patent No. 6900016
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; TITLE OF INVENTION: WITH INFLAMMATORY AUTOIMMUNE DISEASE, M
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; APPLICANT: MCCrae, Keith R.
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Best Local Similarity
Matches 123; Conserv
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Pred. No. 1e-65;
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RESULT 3
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; Sequence 29, Application US/09919039
; Patent No. 6727066
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US-09-919-039-29
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US-09-949-002-476
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APPLICANT: VENTER, J. Craig et al.

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH INFLAMMATORY AUTOIMMUNE DISEASE, METHODS OF DETECTION

TITLE OF INVENTION: AND USES THEREOF

FILE REPERENCE: CL000790

CURRENT APPLICATION NUMBER: US/09/949,002

CURRENT PILING DATE: 2000-01-28

PRIOR FILING DATE: 2000-09-08

PRIOR APPLICATION NUMBER: 60/231,401

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 10823

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 480

LENGTH: 362

TYPE: PRI

TYPE: PR
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Best Local S
Matches 123
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SEQ ID NO 476
LENGTH: 362
TYPE: PRT
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Best Local Similarity 100.0%; Pred. No. 4.3e-65;
Matches 123; Conservative 0; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/09/949,002
CURRENT FILING DATE: 2000-01-28
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nes 123; Conserv
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100.0%; Pred. No. 4.3e-65;
tive 0; Mismatches 0;
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APPLICANT: KASER, Matthew R.
TITLE OF INVENTION: GENES EXPRESSED IN TREATED HUNGLED FOR APPLICATION NUMBER: US/09/919,039
CURRENT FILING DATE: 2002-09-09
PRIOR APPLICATION NUMBER: 60/222,113
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 401
SOFTWARE: PERL Program
SEQ ID NO 29
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH INFLAMMATORY AUTOIMMUNE DISEASE, METHODS OF DETECTION
TITLE OF INVENTION: WITH INFLAMMATORY AUTOIMMUNE DISEASE, METHODS OF DETECTION
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: CLOO0790
CURRENT APPLICATION NUMBER: US/09/949,002
CURRENT FILING DATE: 2000-01-28
PRIOR APPLICATION NUMBER: 60/231,401
PRIOR PRILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 10823
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 475
LENGTH: 579
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence 475, Application US/09949002
Patent No. 6900016
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Best Local Similarity
Matches 123; Conserv
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Best Local Similarity
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ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
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                                           248
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                                                              AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYFTVNCQP 122
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       LGM 125
                                           AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
                                                                                                                  GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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Pred. No. 5.3e-65;
0; Mismatches 0;
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US-08-193-114B-1
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PRIOR FILING DATE: 2000-01-28
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 10823
SOFTWARE: FASUSEQ for Windows Version 4.0
SEQ ID NO 481
LENGTH: 579
TYPE: 779
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH INFLAMMATORY AUTOIMMUNE DISEASE, MITTILE OF INVENTION: AND USES THEREOF
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Best Local (
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CURRENT APPLICATION NUMBER: US/09/949,002
CURRENT FILING DATE: 2000-01-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Jiang, Yongping
TITLE OF INVENTION: Modulation of Blood
TITLE OF INVENTION: Pressure and Inhibition of Platelet Activation
TITLE OF INVENTION: with Kininogen Fragment
NUMBER OF SEQUENCES: 2
                                                APPLICATION NUMBER: US/08/193,114B FILING DATE: 9 February 1994 CLASSIFICATION: 514 PRIOR APPLICATION DATA:
                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720
COMPUTER: IBM PS/2
                                                                                                                                    SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                         STREET: 1800 Two Penn Center Plaza
CITY: Philadelphia
STATE: Pennsylvania
COUNTRY: U.S.A.
                                                                                                                                                                                OPERATING SYSTEM:
APPLICATION NUMBER: U.S. Application
APPLICATION NUMBER: Serial No. 5472945 07/744,545
                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Seidel, Gonda, Lavorgna ADDRESSEE: Monaco, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           308 LGM 310
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100.0%; Pr
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Pred. No.
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PCT-US92-06809-1
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Best Local Similarity
                                                                                                                               CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U.S. Application
APPLICATION NUMBER: Serial No. 744,545
PILING DATE: 13 August 1991
ATTORNEY/AGENT INFORMATION:
NAME: MONACO, DANIEL A.
REGISTRATION NUMBER: 30,480
REFERENCE/DOCKET NUMBER: 6056-137
TELEPHONE: (215) 568-8383
TELEPHONE: (215) 568-5549
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INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acids
                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
COMPUTER: IBM BS/2
OPERATING SYSTEM: MS-DO:
SOFTWARE: WordPerfect 5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/
FILING DATE: 19910813
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REGISTRATION NUMBER: 30,480
REFERENCE/DOCKET NUMBER: 6056
TELECOMMUNICATION: INFORMATION:
TELEPHONE: (215) 568-8383
TELEPAX: (215) 568-5549
                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acid
TYPE: AMINO ACID
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CORRESPONDENCE ADDRESS:
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                    TOPOLOGY:
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                                                         117 amino acids
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|: Mcdulation of Blood
|: Pressure by Altering Bradykinin Levels
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100.0%; Pred. No. 4.5e-59;
htive 0; Mismatches 0;
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Best Local Similarity
                                                                                                                                                  Query Match
Best Local :
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                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US
FILING DATE: Filed Her
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: NOVEL HUMAN CYSTATIN-LIKE TITLE OF INVENTION: PROTEIN
                                                                                                                                                                                                                                  IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: DOS
SOFTWARE: FABISEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                              LENGTH: 178 amino
TYPE: amino acid
STRANDEDNESS: sir
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                                                                                                                                                                                                                    CLONE: 30443
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                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
114 VEIGRTTCKKNQHLRL-DDCDFQTNHTLKQTLSCYSEVWVVPW----VPALRGACSPLSL 168
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                                 71
                                                                                               11 TKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKYFID 70
                                                                                                                               h 25.2%; Score 169; DB 1; Similarity 32.5%; Pred. No. 7.2e-11; 39; Conservative 22; Mismatches 49
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                                                              SRVKPGFPKTIKTNDPGVLQAARYSVEKFNNCTNDMFLFKESRITRALVQIVKGLKYMLE 113
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               FVARETTCSKESNEELTESCE---TKKLGQSLDCNAEVYVVPWEKKIYPTVN--CQPLGM 125
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                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IBM Compatible
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                                                                                                                                                                DB 1; Length 178;
                                                                                                                                 49;
                                                                                                                                  Indels
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                                                                                                                                                                                                         RESULT 11
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                                    Sequence 17, Application US/09461061
Patent No. 6689931
GENERAL INFORMATION:
APPLICANT: McCrae, Keith R.
APPLICANT: McCrae, Keith R.
AITLE OF INVENTION: Inhibition of Angiogenesis By High Molecular Weight
TITLE OF INVENTION: Kininogen Domain 3 Peptide Analogs
FILE REFERENCE: 6056-260 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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CURRENT APPLICATION NUMBER: US/09/461,061
CURRENT FILING DATE: 1999-12-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
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APPLICANT: Bandman, Olga
APPLICANT: Goli, Surya K.
APPLICANT: Goli, Surya K.
OVEL HUMAN CYSTATIN-LIKE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
LENGTH: 178 amino aci
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ATTORNEY/AGENT INFORMATION
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 3174 POI
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Billings, Lucy J
REGISTRATION NUMBER: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
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32.5%; Pred. No. 7.2e-11;
vative 22; Mismatches 49
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EARLIER APPLICATION NUMBER: 60/112,427 EARLIER FILING DATE: 1998-12-16

168

NUMBER OF SEQ ID NOS:

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                                                                                                                                                                                     ; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-832-535-2
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Matches
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Matches
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Patent No. 5919658
GENERAL INFORMATION:
                                                                                                                                       Query Match
Best Local (
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LENGTH: 32
TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/832,535 FILING DATE: 03-APR-1997 CLASSIFICATION: 435
                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS LENGTH: 145 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                              ATTORNBY/AGENT INFORMATION:
NAMES: KIMBALL, PAUL C.
REGISTRATION NUMBER: 34,610
REFERENCE/DOCKET NUMBER: PF:
                                                                                                                                                                                                                                                                                                                 TELEPHONE: (201) 994-1700
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                             71 FVARETTCSKESNEELTESCE---TKKLGOSLDCNAEVYVVPWEKKI-YPTVNC 120
                                                                                           11 TKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKYFID 70
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92 VEIGRTTCKKNQHLRL-DDCDFQTNHTLKQTLSCYSEVWVVPWLQHFEVPVLRC 144
                                                                                                                                         Similarity
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                                                              SRVKPGFPKTIKTNDPGVLQAARYSVEKFNNCTNDMFLFKESRITRALVQIVKGLKYMLE
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                                                                                             HUMAN CYSTATIN F
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96.9%;
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                                                                                                                        Score 165.5; DB 1; Length
Pred. No. 1.3e-10;
22; Mismatches 50; Indels
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Pred. No. 1.6e-11;
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                                                                                                                        Gaps
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Patent No.

91 70

GENERAL INFORMATION:

APPLICANT:

Li, Haodong Yu, Guo-Liang Gentz, Reiner

APPLICANT:

APPLICANT: Ni, Jian
TITLE OF INVENTION: Cystatin
NUMBER OF SEQUENCES: 17 CORRESPONDENCE ADDRESS:

STATE: M CITY: Rockville

3

STREET:

9410 Key West Avenue

ADDRESSEE:

Human

Genome Sciences,

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                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear; MOLECULE TYPE: protein US-09-019-485-2
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US-09-019-485-2
                    US-09-019-485-3
Sequence 3,
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Best Local Similarity
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APPLICANT: Li, Ha
APPLICANT: Yu, Gu
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM FOCOMPACTION
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Benson, Robert H.
REGISTRATION NUMBER: 30,446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
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                                                                                         92 VEIGRTTCKKNOHLRL-DDCDFQTNHTLKOTLSCYSEVWVVPWLQHFEVPVLRC 144
                                                                                                                         71 FVARETTCSKESNEELTESCE---TKKLGQSLDCNAEVYVVPWEKKI-YPTVNC 120
                                                                                                                                                               32 SRVKPGFPKTIKTNDPGVLQAARYSVEKFNNCTNDMFLFKESRITRALVQIVKGLKYMLE
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 Application US/09019485
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Ni, Jian
                                                                                                                                                                                                                                  24.7%; Score 165.5; DB 2; ilarity 32.5%; Pred. No. 1.3e-10; Conservative 22; Mismatches 50;
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                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                    5
                                                                                                                                                                                                                                    Gaps
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ZIP: 20850 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

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APPLICANT: HOLIOWAY, James L.

APPLICANT: Feldhaus, Andrew

TITLE OF INVENTION: TESTIS SPECIFIC CYSTATIN-LIKE PROTEIN CYSTATIN T

FILE REPERBUGE: 98-72

CURRENT APPLICATION NUMBER: US/09/431,480

CURRENT FILING DATE: 1999-11-01

EARLIER APPLICATION NUMBER: 60/109,217

EARLIER APPLICATION NUMBER: 60/156,382

EARLIER APPLICATION NUMBER: 60/156,382

EARLIER FILING DATE: 1999-09-28

NUMBER OF SEQ ID NOS: 22

SOFTWARE: FastSEQ for Windows Version 3.0

SEGUID NO 9

LENGTH: 145

TYPE: PRT

ORGANISN: Homo sapiens

US-09-431-480-9
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; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-09-019-485-3
  밁
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US-09-431-480-9
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                                                                                                                                                                                  Query Match 24.7%; Score 165.5; DB 2; Length 145; Best Local Similarity 32.5%; Pred. No. 1.3e-10; Matches 37; Conservative 22; Mismatches 50; Indels 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9, Application US/09431480 Patent No. 6235708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 3013098439
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Benson, Robert H.
REGISTRATION NUMBER: 30,446
REFERENCE/DOCKET NUMBER: PF265P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 3013098504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/019,485
FILING DATE:
CLASSIFICATION:
92 VEIGRTTCKKNOHLRL-DDCDFQTNHTLKOTLSCYSEVWVVPWLQHFEVPVLRC 144
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                                                                                                                             11 TKICVGCPRDIPTNSPELEETLTHTITKLNABNNATFYFKIDNVKKARVQVVAGKKYFID 70
                                                                                           32 SRVKPGFPKTIKTNDPGVLQAARYSVEKFNNCTNDMFLFKESRITRALVQIVKGLKYMLE 91
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Search completed: February Job time : 48 secs 8, 2006, 03:18:49

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Result
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Maximum Match 100%
Listing first 45 summaries
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Match
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
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291.780 Million cell updates/sec
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    GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration Ltd.
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US-10-661-784-2
US-10-661-784-1
US-10-661-784-1
US-10-693-315-104
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US-10-637-313-20
US-10-637-313-24
US-10-637-313-24
US-10-637-313-24
US-10-637-315-105
US-10-637-315-105
US-10-637-313-14
US-10-637-313-15
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Sequence
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2, Appli
10, Appli
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104, App
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82, Appl
20, Appl
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1179, Ap
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17, Appl
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and

CURRENT APPLICATION NUMBER: US/10/661,784

and

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Sequence 102, Application US/10893315
Publication No. US20050147987A1
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GETTITLE OF INVENTION: WITH TYPE II DIABETES AND TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000786
CURRENT APPLICATION NUMBER: US/10/893,315
CURRENT APPLICATION NUMBER: US/10/893,315
        RESULT 4
US-10-893-315-104
US-10-893-315-104
; Sequence 104, Application US/10893315
; Publication No. US20050147987A1
; GENERAL INFORMATION:
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PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin version 3.2
SEQ ID NO 1
LENGTH: 123
TYPE: PRT
ORGANISM: Homo sapiens
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; ORGANISM: Human
US-10-893-315-102
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US-10-893-315-102
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SEQ ID NO 102
LENGTH: 362
TYPE: PRT
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Best Local (
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Best Local Similarity
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PRIOR APPLICATION NUMBER: 60/231,397
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 2172
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Craig et al
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100.0%; Pred. No. 2.9e-61;
tive 0; Mismatches 0;
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; LENGTH: 362
; TYPE: PRT
; ORGANISM: Human
US-10-893-315-104
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US-10-162-335-82
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Best Local Similarity
Matches 123; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH TYPE II DIABETES AND OBESITY, METHODS OF DETECTION AND
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CLOO0786
CURRENT APPLICATION NUMBER: US/10/893,315
CURRENT FILING DATE: 2004-07-19
PRIOR APPLICATION NUMBER: 60/231,397
PRIOR PPLICATION NUMBER: 60/231,397
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 2172
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APPLICANT:
APPLICANT:
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PRIOR APPLICATION NUMBER: 60/295,607
PRIOR FILLING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: 60/295,661
PRIOR FILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: 60/296,404
PRIOR FILING DATE: 2001-06-06
                                                                                                             APPLICANT: Zerhusen, Bryan D.
TITLE OF INVENTION: Therapeutic Polypeptides,
FILE REFERENCE: 21402-377 B
CURRENT APPLICATION NUMBER: US/10/162,335
CURRENT FILING DATE: 2002-10-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Anderson, David W. APPLICANT: Baumgartner, Jaso
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Millet, Isabelle
Padigaru. M....
                                                                                                                                                                                                         Stone, David J.
Spytek, Kimberly A.
Vernet, Corine A. M.
Voss, Edward Z.
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Pena, Carol E. A.
Rastelli, Luca
Shimkets, Richard A.
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Gangolli,
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Gorman, Linda
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t, Tord
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, Stacie J.
r, Shlomit R.
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                                                                                                                                                                          Nucleic Acids Encoding Same, and Methor
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RESULT 6
US-10-637-313-32
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                                                                              CURRENT APPLICATION NUMBER: US/10/637,313
CURRENT PILING DATE: 2003-08-08
PRIOR APPLICATION NUMBER: 10/16235
PRIOR PILLING DATE: 2002-06-03
PRIOR APPLICATION NUMBER: 60/295607
PRIOR FILLING DATE: 2001-06-04
PRIOR PILLING DATE: 2001-06-04
PRIOR PILLING DATE: 2001-06-04
PRIOR PILLING DATE: 2001-06-04
PRIOR PILLING DATE: 2001-06-06
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Best Local Similarity
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LENGTH: 390
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APPLICANT: Uriel
APPLICANT: Li Li
                                                                                                                                                                                                                                                                                        APPLICANT: Corine Vernet

APPLICANT: Meera Patturajan

TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME,

FILE REFERENCE: 21402-593B

TILE REFERENCE: 21402-593B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OR APPLICATION NUMBER: 60/297,567
OR FILING DATE: 2001-06-12
OR APPLICATION NUMBER: 60/298,285
OR FILING DATE: 2001-06-14
OR APPLICATION NUMBER: 60/298,556
OR FILING DATE: 2001-06-15
OR APPLICATION NUMBER: 60/299,949
OR APPLICATION NUMBER: 60/290,949
OR APPLICATION NUMBER: 60/300,883
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APPLICATION NUMBER: 60/296418
FILING DATE: 2001-06-06
APPLICATION NUMBER: 60/298285
FILING DATE: 2001-06-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   63
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Nikolai Khramtsov
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No. US20040229779A1
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US-10-637-313-32
APPLICANT: Zerhusen, Bryan D.
TITLE OF INVENTION: Therapeutic Polypeptides, Nu
FILE REFERENCE: 21402-377 B
CURRENT APPLICATION NUMBER: US/10/162,335
CURRENT FILING DATE: 2002-10-01
PRIOR APPLICATION NUMBER: 60/295,607
PRIOR APPLICATION NUMBER: 60/295,661
PRIOR APPLICATION NUMBER: 60/296,404
PRIOR APPLICATION NUMBER: 60/296,404
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SEQ ID NO 32
LENGTH: 390
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Best Local Similarity
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PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/299949
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/300883
PRIOR FILING DATE: 2001-06-26
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PRIOR FILING DATE: 2001-06-28
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Spytek, Kimberly A.
Vernet, Corine A. M.
Voss, Edward Z.
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Shimkets, Richard A.
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Malyankar, Uriel
Millet, Isabelle
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Hjalt, Tord
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Gorman, Linda
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Pena, Carol
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                                                                                                                                                                                                                                                                                             Meera
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    See File Wrapper or PALM

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             APPLICANT: Corine Vernet
APPLICANT: Meera Patturajan
TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NU
FILE REFERENCE: 21402-593B
CURRENT APPLICATION NUMBER: US/10/637,313
CURRENT FILING DATE: 2003-08-08
PRIOR APPLICATION NUMBER: 10/162335
PRIOR FILING DATE: 2002-06-03
PRIOR APPLICATION NUMBER: 60/295607
PRIOR FILING DATE: 2001-06-04
PRIOR PILING DATE: 2001-06-04
PRIOR FILING DATE: 2001-06-04
PRIOR FILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: 60/2956404
PRIOR APPLICATION NUMBER: 60/296418
PRIOR APPLICATION NUMBER: 60/296418
PRIOR FILING DATE: 2001-06-06
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US-10-637-313-20
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Best Local Similarity
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Remaining Prior Application data removed - See File Wrapper or PALM
NUMBER OF SEQ ID NOS: 201
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APPLICANT: Uriel Malyankar
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OR APPLICATION NUMBER: 60/298,556
OR FILING DATE: 2001-06-15
(OR APPLICATION NUMBER: 60/299,949
(OR APPLICATION NUMBER: 60/299,949
(OR APPLICATION NUMBER: 60/299,949
(OR APPLICATION NUMBER: 60/290,949)
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RAPPLICATION NUMBER: 60/296,418

OR FILING DATE: 2001-06-06

OR APPLICATION NUMBER: 60/297,414

OR FILING DATE: 2001-06-11

OR PILING DATE: 2001-06-12

OR FILING DATE: 2001-06-12
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Ferenc Boldog
Charles Miller
Nikolai Khramtsov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/10637313
No. US20040229779A1
                                                                                                                                                                                                                                                                                                                                                                                                 Mei Zhong
                                                                                                                                                                                                                                                                                                                                                                              Muralidhara Padigaru
                                                                                                                                                                                                                                                                                                                                                                                                                    Sasha Guo
                                                                                                                                                                                                                                                                                                                                                                                                                                   David Anderson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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373
                                   123
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; SEQ ID NO 20
; LENGTH: 398
; TYPE: PRT
; ORCANISM: Homo sapiens
US-10-637-313-20
                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID
US-09-919-039-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-919-039-29
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PRIOR APPLICATION NUMBER: 60/29856
PRIOR APPLICATION NUMBER: 60/299949
PRIOR APPLICATION NUMBER: 60/299949
PRIOR APPLICATION NUMBER: 60/30083
PRIOR FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: 60/301550
PRIOR APPLICATION NUMBER: 60/301550
PRIOR APPLICATION NUMBER: 60/301550
PRIOR APPLICATION NUMBER: 60/301550
PRIOR FILING DATE: 2001-06-26
PRIOR HILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: 60/301550
PRIOR DATE: 2001-06-26
PRIOR                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/919,039
CURRENT FILING DATE: 2002-09-09
PRIOR APPLICATION NUMBER: 60/222,113
PRIOR FILING DATE: 2000-07-28
RUMBER OF SEQ ID NOS: 401
SOFTWARE: PERL Program
SEQ ID NO 29
LENGTH: 427
TYPET
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                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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TITLE OF INVENTION: GENES EXPRESSED IN TREATED HUMAN C3A LIVER CELL CULTURES FILE REFERENCE: PA-0035 US
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Local Similarity 100.0%; Pred. No. 3.2e-61;
Les 123; Conservative 0; Mismatches 0;
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                                                                                             AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYFTVNCQP 122
                                                                                                                                                                                                         GKDFVQPPTKI CVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKI DNVKKARVQVV
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LGM 125
                                                         AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
                                                                                                                                                                        GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                 Length 427;
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**LGM 375** 

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LENGTH: 427
; TYPE: PRT
; ORGANISM: Homo s
US-10-637-313-24
RESULT 11
US-10-741-600-1179
; Sequence 1179, Application US/10741600
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CURRENT APPLICATION NUMBER: US/10/637,313
CURRENT FILING DATE: 2003-08-08
PRIOR APPLICATION NUMBER: 10/162335
PRIOR APPLICATION NUMBER: 60/295607
PRIOR APPLICATION NUMBER: 60/295661
PRIOR PILING DATE: 2001-06-04
PRIOR PILING DATE: 2001-06-04
PRIOR PILING DATE: 2001-06-06
PRIOR PILLING DATE: 2001-06-06
PRIOR PILLING DATE: 2001-06-06
PRIOR PILLING DATE: 2001-06-06
PRIOR PILING DATE: 2001-06-06
PRIOR PILING DATE: 2001-06-06
PRIOR PILING DATE: 2001-06-06
PRIOR PILING DATE: 2001-06-14
PRIOR PILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: 60/29828
PRIOR APPLICATION NUMBER: 60/29828
PRIOR APPLICATION NUMBER: 60/29828
PRIOR PILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: 60/29828
PRIOR APPLICATION NUMBER: 60/29849
PRIOR APPLICATION NUMBER: 60/29949
PRIOR APPLICATION NUMBER: 60/20949
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US-10-637-313-24
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APPLICANT: Uriel M
APPLICANT: Li Li
APPLICANT: David A
APPLICANT: Sasha C
APPLICANT: Mei Zhc
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Best Local Similarity
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Publication No. US20040229779A1
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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PRIOR FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: 60/301550
PRIOR FILING DATE: 2001-06-28
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TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME,
                                                                                                                                                             123
                                                                                                                     373
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                                                                                                                                                                                                                                                                                                 253 GKDFVQPPTKICVGCPRDIPTNSPELBETLTHTITKLNAENNATFYFKIDNVKKARVQVV 312
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                                                                                                                                                                LGM 125
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Nikolai Khramtsov
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Ferenc Boldog
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                                                                                                                                                                                                                                                                                                                                                                                        98.5%; Score 660; DB 5; 1 (larity 100.0%; Pred. No. 3.5e-61; Conservative 0; Mismatches 0;
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GENERAL INFORMATION:

APPLICANT: CARGILL, Michele et al.

TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH

TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF DETECT

FILE REFERENCE: CL001499

CURRENT APPLICATION NUMBER: US/10/741,600

CURRENT FILING DATE: 2003-12-22

NUMBER OF SEQ ID NOS: 73997

SOFTMARE: FastSEQ for Windows Version 4.0

SEQ ID NO 1181

LENGTH: 427

TYPE: PRT
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US-10-741-600-1181
; Sequence 1181, Application US/10741600
; Publication No. US20050026169A1
; Publication...
RESULT 13
US-10-893-315-101
; Sequence 101, Application US/10893315
; Publication No. US20050147987A1
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-741-600-1179
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TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED I
TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF THE REFERENCE: CL001499
CURRENT APPLICATION NUMBER: US/10/741,600
CURRENT FILING DATE: 2003-12-22
NUMBER OF SEQ ID NOS: 73997
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 98.5%; Score 660; DB 5; I Best Local Similarity 100.0%; Pred. No. 3.5e-61; Matches 123; Conservative 0; Mismatches 0;
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                                                                                                                                                      375
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; PRIOR APPLICATION NUMBER: 60/231,397
; PRIOR FILING DATE: 2004-07-19
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 2172
; SOPTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 105
; SEQ ID NO 105
; LENGTH: 579
; ORGANISM: Human
US-10-893-115-107
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TITLE OF INVENTION: POLYMORPHISMS IN KNO!
TITLE OF INVENTION: WITH TYPE II DIABET!
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000786
CURRENT APPLICATION NUMBER: US/10/893,31:
CURRENT FILING DATE: 2004-07-19
PRIOR APPLICATION NUMBER: 60/231,397
PRIOR APPLICATION NUMBER: 60/231,397
PRIOR FILING DATE: 2000-09-08
INMBER OF SEG ID NOS: 2172
SOFTWARE: FastSEQ for Windows Version 4.0
SEG ID NO 101
LENGTH: 579
TYPE: PRT
TYPE: PRT
TYPE: PRT
TYPE: PRT
TYPE: PRT
TOGRANISM: Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 105, Application US/10893315 Publication No. US20050147987A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 123; Conserv
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN
TITLE OF INVENTION: WITH TYPE II DIABETES
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CLOOO786
CURRENT PEPLICATION UNDERE: US/10/893,315
CURRENT FILING DATE: 2004-07-19
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Similarity 100.0%; Pred. No. 5.2e-61;
                                    LGM 125
                                                                         AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
                                                                                                  AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 122
                                                                                                                                                      GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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LGM 310
                                                                                                                                                                                                                                98.5%; Score 660; DB 5; I ilarity 100.0%; Pred. No. 5.2e-61; Conservative 0; Mismatches 0;
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US-10-162-335-72
Sequence 72, Application US/10162335
Publication No. US20040009480A1
                                                                                                                                                                                                                               US-10-162-335-72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/296,404
PRIOR FILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: 60/296,418
PRIOR PILLING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: 60/297,414
PRIOR FILING DATE: 2001-06-11
PRIOR APPLICATION NUMBER: 60/297,567
PRIOR FILING DATE: 2001-06-12
PRIOR APPLICATION NUMBER: 60/298,285
                                                                                                                                                                                                                                                                                                        Remaining Prior Application NUMBER OF SEQ ID NOS: 201 SEQ ID NO 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                               Matches
                                                                                                                                                                                  Query Match
                                                                                                                                                                  Best Local Similarity
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TITLE OF INVENTION: Therapeutic Polypeptides, Nucleic Acids Encoding Same, and Method PILE REFERENCE: 21402-377 B
CURRENT APPLICATION NUMBER: US/10/162,335
CURRENT FILING DATE: 2002-10-01
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TYPE: PRT
ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2002-10-01
PRIOR APPLICATION NUMBER: 60/295,607
PRIOR FILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: 60/295,661
PRIOR FILING DATE: 2001-06-04
                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/300,883 PRIOR FILING DATE: 2001-06-26
                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/298,556
PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/299,949
PRIOR FILING DATE: 2001-06-21
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AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYFTVNCQP
                                                             GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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Pena, Carol E. A.
Rastelli, Luca
Shimkets, Richard A.
                                                                                                       GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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Gerlach, Valerie
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Boldog, Ferenc L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Spytek, Kimberly A. Vernet, Corine A. M. Voss, Edward Z.
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Malyankar, Uriel M.
Millet, Isabelle
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Hjalt, Tord
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Casman, Stacie J.
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                                                                                                                                          98.5%; Score 660; DB 4; ilarity 100.0%; Pred. No. 5.6e-61; Conservative 0; Mismatches 0;
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		Search completed: February ( Job time : 179 sécs	284 AGKKYFIDFVARETT 123 LGM 125     344 LGM 346
		8, 2006, 03:21:54	rcskesnebltescetkklg
			AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 343 LGM 125 LGM 346
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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  Score
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seq length: 2000000000
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2: /cgn2_6/ptodata/2/pubpaa/USOB_NEW_PUB.pep:*
3: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
4: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
5: /cgn2_6/ptodata/2/pubpaa/USO3_NEW_PUB.pep:*
6: /cgn2_6/ptodata/2/pubpaa/USO3_NEW_PUB.pep:*
7: /cgn2_6/ptodata/2/pubpaa/USI1_NEW_PUB.pep:*
8: /cgn2_6/ptodata/2/pubpaa/USO3_NEW_PUB.pep:*
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Match
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US-10-967-093-8
US-10-967-093-10
US-10-967-093-4
US-10-967-093-4
US-10-967-093-2
US-10-988-962-7
US-10-888-962-8
US-11-19-041-23
US-11-19-041-34-955
US-10-877-346-40
US-10-877-346-40
US-11-19-041-138
US-11-19-041-138
US-11-19-041-138
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Sequence 8, Appli
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16, Appli
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18, Appli
19, Appli
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61.5 9.2 384 7 US-11-075-351-12 61.5 9.2 178 6 US-10-943-89-29 61.5 9.2 1178 7 US-11-044-89-29 61.5 9.2 1286 6 US-10-877-346-38 61.5 9.2 2053 6 US-10-877-346-11 61.5 9.2 2066 6 US-10-877-346-11 61.5 9.2 2066 6 US-10-877-346-11 61 9.1 227 6 US-10-485-517-210 61 9.1 326 6 US-10-485-517-241 61 9.1 3106 6 US-10-485-517-241 61 9.1 3106 6 US-10-485-517-241 60.5 9.0 334 7 US-11-016-564-1 60.5 9.0 407 7 US-11-016-564-7 60.5 9.0 407 7 US-11-016-564-7 60.5 9.0 407 7 US-11-016-564-7 60.5 9.0 603 7 US-11-183-136-2 60 9.0 651 6 US-10-851-2341-366 59.5 8.9 193 5 US-09-978-360A-558 59.8 8.9 248 7 US-11-054-515-1386 59.8 8.9 248 7 US-11-054-515-1386	9.2 384 7 US-11-075-351-12 9.2 1628 (10-71)-626-28 9.2 1178 7 US-11-044-899-28 9.2 1178 6 US-10-877-346-38 9.2 1286 6 US-10-877-346-38 9.2 2056 6 US-10-877-346-19 9.2 2066 6 US-10-485-517-24 9.1 1306 6 US-10-485-517-24 9.1 1306 6 US-10-486-517-24 9.1 1306 6 US-10-486-517-24 9.1 1306 6 US-10-486-517-24 9.1 1306 6 US-10-486-517-24 9.1 1306 6 US-10-486-54-1 9.0 391 7 US-11-016-564-1 9.0 603 7 US-11-1016-564-1 9.0 603 7 US-11-183-136-2 9.0 603 7 US-11-183-136-2 9.0 651 6 US-10-995-548-3 9.0 651 6 US-10-995-561-95 8.9 481 6 US-10-995-561-95 8.8 948 7 US-11-054-515-19	45	44	43	42	41	40	39	38	37	36	35	ω 4	ω u	32	31	30	29	28	27	
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US-11-075-351-12 US-11-074-899-25 US-11-044-899-25 US-10-877-346-38 US-10-877-346-31 US-10-877-346-11 US-10-485-517-21 US-10-485-517-54 US-11-016-564-1 US-11-016-564-1 US-11-106-564-1 US-11-106-564-1 US-11-108-564-1 US-11-108-564-1 US-11-108-564-1 US-11-016-564-1 US-11-052-554A-3 US-11-052-554A-3 US-10-995-561-95 US-11-052-554A-3 US-10-995-561-95	US-11-075-351-12 US-10-793-626-2844 US-10-494-899-29 US-11-044-899-29 US-11-044-899-29 US-10-877-346-38 US-10-877-346-11 US-10-877-346-11 US-10-485-517-210 US-10-485-517-210 US-10-485-517-241 US-10-485-517-241 US-10-485-517-241 US-10-467-657-4406 US-11-016-564-1 US-11-016-564-1 US-11-016-564-7 US-11-016-564-7 US-11-016-564-7 US-11-016-564-7 US-11-016-564-7 US-11-016-564-7 US-11-018-584-300 US-11-018-584-300 US-11-018-584-300 US-11-018-584-300 US-11-018-584-300 US-11-018-584-300 US-11-018-584-300 US-11-052-584-300 US-11-052-584-300 US-10-821-234-13666 US-10-821-234-13666 US-10-985-561-959 US-11-054-515-1386	932	248	481	193	651	267	603	407	391	334	1306	227	227	2066	2053	1286	1178	628	384	
-11-075-351-12 -10-075-352-12 -11-044-899-29 -10-877-346-38 -10-877-346-9 -10-877-346-9 -10-485-517-21 -10-485-517-24 -10-485-517-54 -11-016-564-1 -11-016-564-7 -11-016-564-7 -11-183-136-2 -11-018-254A-7 -11-052-254A-7 -11-052-254A-7 -11-052-254A-7 -11-052-254A-7	-11-075-351-12 -10-793-26-2844 -10-47-39-29 -10-47-346-38 -10-877-346-38 -10-877-346-11 -10-877-346-9 -10-877-346-9 -10-485-517-210 -10-485-517-241 -10-467-557-5406 -11-016-564-1 -11-016-564-7 -11-0	7	7	0	v	σ	7	7	7	7	7	σ	σ	σ	o	σ	σ	7	σ	7	
		US-11-057-058-59	US-11-054-515-1386	US-10-995-561-959	US-09-978-360A-558	US-10-821-234-1666	-11-052-554A-	US-11-183-136-2	-11	Ŀ	US-11-016-564-1	US-10-467-657-5406	US-10-485-517-241	US-10-485-517-210	US-10-877-346-9	US-10-877-346-11	-10-877-346-3		-10-793-626-28	-11-075-351	

## ALIGNMENTS

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RESULT 1
US-10-689-742-197
; Sequence 197, Application US/10689742
; Sequence 197, Application US/10689742
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US-10-967-093-8
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Sequence 8, Application US/10967093
Publication No. US20050267021A1
GENERAL INFORMATION:
APPLICANT: Schiemann, William P.
TITLE OF INVENTION: Cystatin C as
FILE REFERENCE: 2879-104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Jacobs, Kenneth
APPLICANT: McCoy, John M
APPLICANT: LaVallie, Edward R
APPLICANT: Racie, Lisa A
APPLICANT: Evans, Cheryl
APPLICANT: Mcrberg, David
APPLICANT: Treacy, Maurice
                                                                                                                                                                                                                                                                                                                                                                                                    Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: 00766.000091.10
FULE REPERENCE: 00766.000091.10
CURRENT APPLICATION NUMBER: US/10/689,742
CURRENT FILING DATE: 2003-10-22
PRIOR APPLICATION NUMBER: 09/746,783
PRIOR APPLICATION NUMBER: 09/746,783
PRIOR FILING DATE: 2000-12-21
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                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
mes 37; Conserv
                                                                                                                                                                                                                        114 VEIGRTTCKKNQHLRL-DDCDFQTNHTLKQTLSCYSEVWVVPWLQHFEVPVLRC 166
                                                                                                                                                                                                                                                  71 FVARETTCSKESNEELTESCE---TKKLGQSLDCNAEVYVVPWEKKI-YPTVNC 120
                                                                                                                                                                                                                                                                                                           54 SRVKPGFPKTIKTNDPGVLQAARYSVEKENNCTNDMFLFKESRITRALVQIVKGLKYMLE
                                                                                                                                                                                                                                                                                                                                                  11 TKICYGCPRDIPTNSPELBETLTHTITKLNAENNATFYFKIDNVKKARYQVVAGKKYFID
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                                                                                                                                                                                                                                                                                                                                                                                             24.7%; Score 165.5; DB 6; 32.5%; Pred. No. 1.6e-10; tive 22; Mismatches 50;
                         an Antagonist of
                           TGF-B
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                             Methods
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                             Related
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Sequence 6, Application US/10967093
Publication No. US20050267021A1
GENERAL INFORMATION:
APPLICANT: Schiemann, William P.
TITLE OF INVENTION: Cystatin C as an Antagonist of FILE REFERENCE: 2879-104
CURRENT APPLICATION NUMBER: US/10/967,093
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: 60/511,794
PRIOR APPLICATION NUMBER: 60/511,794
PRIOR APPLICATION NUMBER: 60/511,794
PRIOR FILING DATE: 2003-10-15
NUMBER OF SEQ ID NOS: 10
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Matches
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SOFTWARE: PatentIn version 3.1 SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local Similarity
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-10-967-093-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Schiemann, William P.
TITLE OF INVENTION: Cystatin C as an Antagonist of TGF-B and Methods Related Thereto FILE REFERENCE: 2879-104
CURRENT SPELICATION NUMBER: US/10/967,093
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: 60/511,794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/511,794 PRIOR FILING DATE: 2003-10-15 NUMBER OF SEQ ID NOS: 10
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NUMBER OF SEQ ID NOS:
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CURRENT FILING DATE: 2004-10-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Macaca mulatta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 106 QANLDSCPFHNQPHLKREKL----CSFQVYVVPWMN----TINLVKFSCQ 147
                                                                                                                                                                                                                                                                                                                                                                    89
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                                                                                                                                                                                                                                                                                                                                                                                                          68 FIDFVARETTCSKESNEELTESC---ETKKLGQSLDCNAEVYVVPWE 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                     31 KPPR--LVGGPMDASVEEEGVRRALDFAVSEYNKASNDMYHSRALQVVRARKQIVAGVNY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 QPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKY 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                  FLDVELGRTTCTK--TQPNLDNCPFHEQPHLKRKAFCSFQIYTVPWQ 133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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29.0%; Pred. No. 3e-0
ative 21; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            48;
                                                                                                                                                                         of TGF-B
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                                                                                                                                                                         and Methods Related Thereto
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Sequence 425, Application US/09978360A

| Publication No. US20060009633A9
| GENERAL INFORMATION:
| APPLICANT: Edwards, Jean-Baptiste Dumas Milne
| APPLICANT: Duclert, Aymeric
| APPLICANT: Duclert, Severin
| APPLICANT: Jobert, Severin
| APPLICANT: Jobert, Severin
| APPLICANT: Clused, Catherine
| TITLE OF INVENTION: Complementary DNA's Encoding Proteins with Signal Peptides
| FILE REFERENCE: 56.US4.CIP
| CURRENT FILING DATE: 2001-10-15
| PRIOR APPLICATION NUMBER: US 60/066,677
| PRIOR APPLICATION NUMBER: US 60/066,677
| PRIOR APPLICATION NUMBER: US 60/069,957
| PRIOR APPLICATION NUMBER: US 60/069,957
| PRIOR APPLICATION NUMBER: US 60/074,121
| PRIOR APPLICATION NUMBER: US 60/081,563
| PRIOR APPLICATION NUMBER: US 60/081,563
| PRIOR APPLICATION NUMBER: US 60/081,563
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US-10-967-093-4
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Publication No. US20050267021A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local Similarity
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CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: 60/511,794
PRIOR FILING DATE: 2003-10-15
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Schiemann, William P. TITLE OF INVENTION: Cystatin C as an Antagonist of TGF-B and Methods Related Thereto FILE REFERENCE: 2879-104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 140
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Local Similarity 30.0%;
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Pred. No. 6.5e-07;
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; LOCATION: -15..-1
US-09-978-360A-425
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Matches
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Best Local Similarity
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SEQ ID NO 425
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Publication No. US2005024480001
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PRIOR
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                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 016866-011550US
CURRENT APPLICATION NUMBER: US/10/982,545
CURRENT FILING DATE: 2004-11-06
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PRIOR
                                                                                                                             PRIOR FILING DATE: 2004-05-16 PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2004-07-08
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TITLE OF INVENTION: Biomarkers for Alzheimer's Disease
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APPLICANT: McGuire, James
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                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo Sapiens
          TYPE: PRT
ORGANISM: Homo sapiens
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EATURE:
                                                    ENGTH: 146
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FILING DATE: 1998-12-17
APPLICATION NUMBER: PCT/IB98/02122
FILING DATE: 1998-12-17
APPLICATION NUMBER: US 09/247,155
FILING DATE: 1999-02-09
                                                                                                                                                                                                                                             APPLICATION NUMBER: US 60/546,423
FILING DATE: 2004-02-19
APPLICATION NUMBER: US 60/547,250
FILING DATE: 2004-02-23
                                                                                                                                                                    APPLICATION NUMBER: US 60/572,617 FILING DATE: 2004-05-18
                                                                                                                                                                                                        APPLICATION NUMBER: US (FILING DATE: 2004-04-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/099,273 FILING DATE: -09-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 60/096,116
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                                                                                         PatentIn
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Simonsen, Anja Hvi
Blennow, Kaj
Podust, Vladimir
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Best Local Similarity
Matches 30; Conserv
                                                                                                         Sequence 1237, Application US/10821234 Publication No. US20050255114A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1 SEQ ID NO 2
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             APPLICANT: Andarmani, Susan
APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Schiemann, William P.
TITLE OF INVENTION: Cystatin C as an Antagonist of TGF-B and Methods Related Thereto FILE REFERENCE: 2879-104
CURRENT APPLICATION NUMBER: US/10/967,093
CURRENT FILING DATE: 2004-10-15
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NAME/KBY: SIGNAL
LOCATION: (1)..(26)
OTHER INFORMATION: 6
                                                                      APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 10
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PRIOR FILING DATE: 2003-10-15
FILE REFERENCE: 821A
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TYPE: PRT
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LOCATION: (35)..(146)
OTHER INFORMATION: biomarker peptide M12583.4, truncated Cystatin
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OTHER INFORMATION: biomarker peptide M13391, full-length Cystatin
OTHER INFORMATION: mature peptide
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28.0%; Pro
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Pred. No. 1.3e-06;
1; Mismatches 49;
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                 and Treatment
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                 of Preeclampsia
                                                                                                                                                                                                                                             133
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                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Sequence 8, Application US/10888962
Publication No. US20050266531A1
GENERAL INFORMATION:
APPLICANT: Saint Louis University
APPLICANT: Ray, Ranjit
APPLICANT: Yie-Hwa, Chang
APPLICANT: Ray, Ratna
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US-10-888-962-7
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                                                                                                                                                          US-10-888-962-8
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Best Local S
Matches 25
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CURRENT FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
PRIOR FILING DATE: 2103-04-07
NUMBER OF SEQ ID NOS: 1704
SOFTWARE: pt_SEQ_genes Version 1.0
SEQ ID NO 1237
LENGTH: 97
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Matches
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PRIOR FILING DATE: 2003-07-12
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.2
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 7, Application US/10888962 Publication No. US20050266531A1 GENERAL INFORMATION:
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APPLICANT: Ray, Ranjit
APPLICANT: Yie-Hwa, Chang
APPLICANT: Ray, Ratna
APPLICANT: Basu, Arnab
TITLE OF INVENTION: Compositions and Methods for Inhibiting Liver Stellate Cell
TITLE OF INVENTION: Growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: SLU 03-013 PCT
CURRENT APPLICATION NUMBER: US/10/888,962
CURRENT FILING DATE: 2004-07-09
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 345
TYPE: PRT
ORGANISM: Mus musculus
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                                                                                                                                                                                                                                     192 PLPVSTLVEFVIAATDCTA---KEVTDPAKCNLLAE 224
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Similarity 28.9%;
24; Conservative 19
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25; Conservative
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                                                                                                                                                                                                                                                                            VAGKKYFIDFVARETTCSKESNEELTESCETKKLGQ 97
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                                                                                                                                                                                                                                                                                                                                                                                               18; Mismatches
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Best Local Similarity
Watches 26; Conserve
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TITLE OF INVENTION: Growth
FILE REFERENCE: SLU 03-013 PCT
CURRENT APPLICATION NUMBER: US/10/888,962
CURRENT FILING DATE: 2004-07-09
PRIOR APPLICATION NUMBER: 60/487,126
PRIOR APPLICATION NUMBER: 60/487,126
PRIOR FILING DATE: 2003-07-12
SUPERIOR OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.2
SEQ ID NO 8
LENGTH: 352
LENGTH: 352
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SEQ ID NO 6
LENGTH: 367
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Publication No. US20050266531A1
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Best Local (
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APPLICANT: Ray, Ranjit
APPLICANT: Yie-Hwa, Chang
APPLICANT: Ray, Ratna
APPLICANT: Basu, Ratna
APPLICANT: Basu, Arnab
TITLE OF INVENTION: Growth
TITLE OF INVENTION: Growth
TITLE OF INVENTION: Growth
TITLE REPERENCE: SLU 03-013 PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/888,962
CURRENT FILING DATE: 2004-07-09
PRIOR APPLICATION NUMBER: 60/487,126
PRIOR FILING DATE: 2003-07-12
NUMBER OF SEQ ID NOS: 15
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ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                        LENGTH: 367
TYPE: PRT
ORGANISM: Homo mapiens
                                                                                                                                                                                                                                                                       Match 13.2%; Score 88.5; DB 6; Local Similarity 26.1%; Pred. No. 0.048;
                                                                                                                                                             134 SSPDSAEDVRKYCQDCPLLAPLNDTRVVHAAKAALAAFNAQNNGS-NFQLEEISRAQLVP 192
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247
                                                                              193 LPPSTY-VEFTVSGTDCVAKEATEAAKCNLLAEKQYGFCKATLSEKLG---
                                       109 PWEKKIYPTVNCQP 122
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                                                                                                                      62 VAGKKYFIDFVARETTC-SKESNEE-----LTES----CE---TKKLGQSLDCNAEVYVV 108
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CTVFQTQP-VTSQP
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                                                                                                                                                                                                                                                       Conservative
259
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RESULT 13 US-10-467-962B-12

; Sequence 12,

Application US/10467962B

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ORGANISM: Homo sapiens US-11-137-465-37
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CURRENT APPLICATION NUMBER: US/11/137,465
CURRENT FILING DATE: 2005-05-25
PRIOR APPLICATION NUMBER: US/10/239,663
PRIOR FILING DATE: 2002-09-24
PRIOR APPLICATION NUMBER: PCT/US01/09226
PRIOR FILING DATE: 2001-03-22
PRIOR PILING DATE: 2000-03-24
PRIOR PILING DATE: 2000-03-24
PRIOR PILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: 60/192,668
PRIOR APPLICATION NUMBER: 60/200,166
PRIOR APPLICATION NUMBER: 60/200,166
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US-11-137-465-37
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  Query Match 11.0%;
Best Local Similarity 24.5%;
Matches 23; Conservative 2
                                                                                                                                                                              SEQ ID NO 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 12.2%; Score 82; DB 6 Best Local Similarity 23.5%; Pred. No. 0.57; Matches 32; Conservative 23; Mismatches
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APPLICANT:
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APPLICANT: Murdoch, Paul
APPLICANT: Rizvi, Safia,
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APPLICANT: Klein, Mathieu
TITLE OP INVENTION: Identification of Herbicidally Active Substances
FILE REFERENCE: 2000 857
FILE REFERENCE: 2000 857
CURRENT APPLICATION NUMBER: US/10/467,962B
CURRENT FILING DATE: 2003-08-14
PRIOR APPLICATION NUMBER: PCT/EP02/01466
PRIOR FILING DATE: 2002-02-13
NUMBER: OF SEQ ID NOS: 109
NUMBER: OF SEQ ID NOS: 109
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                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 66
SOFTWARE: FastSEQ for Wi
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TITLE OF INVENTION: NOVEL COMPOUNDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Plésch, Gunnar
APPLICANT: Blau, Astrid
APPLICANT: Blau, Astrid
APPLICANT: Daeschner, Klaus
APPLICANT: Klein, Mathieu
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                                                                                                                                    LENGTH: 13
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 188 IGKLMDMEKASTVDLMRLILSYVVDVASSEE--SKQHNKEIMESSVRSLLSQI----AKM
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Rizvi, Safia, K.
Smith, Randall, F.
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                                                                                                                                                                                              for Windows Version 3.0
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20;
    Score 74; DB
Pred. No. 0.47
20; Mismatches
                        DB 7; Length 138; 0.47;
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    47; Indels
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  Gaps
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Search completed: February Job time : 17 secs

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US-10-888-962-5
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US-10-888-962-5
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                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: SLU 03-013 PCT CURRENT APPLICATION NUMBER: US/10/888,962 CURRENT FILING DATE: 2004-07-09 PRIOR APPLICATION NUMBER: 60/487,126 PRIOR FILING DATE: 2003-07-12 NUMBER OF SEQ ID NOS: 15
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TITLE OF INVENTION: Compositions and Methods
TITLE OF INVENTION: Growth
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APPLICANT: Ray, Ranjit
APPLICANT: Yie-Hwa, Ch
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                                                                                                                                               Local Similarity
les 23; Conserv
 190 ---
                                                                     134 SSPDSAEDVRKLCPDCPLLAPLNDSRVVHAVEVALATFNAESNGS-YLQLVEISRAQ---
                                62 VAGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCN 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          92
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                                                                                                         2 SGKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQV 61
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No. US20050266531A1
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                                                                                                                                               Conservative
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FVPLPVSVSVEFAVAATD-CIAKEVVDPTKCN
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Perfect score:
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80 C	44444 4544 8888	91.5 13.7 88.5 13.2 88.5 13.2 88.5 13.2 88.5 13.2 85 12.7 84.5 12.6	375 2 125 2 134 2 134 2 134 2 367 1 199 2 592 2	A32827 T00752 JC4882 S54828 WOHU S65071 T50516	fetuin precursor - cysteine proteinas cystatin - maize cysteine proteinas alpha-2-HS-glycopr cystatin - field m Ca2+ dependent lip
				ALIGNMENTS	
	RESULT 1 KGHUL1 kininogen,	MW precursor	[valida	ated] - human	
	N;Alternate N;Contains:	names: alpna bradykinin (	-2-thio	l proteinase inhibitor; a); kininogen I; kinino	N;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen N;Contains: bradykinin (kallidin); kininogen I; kininogen II; prokininogen C.Soncies: Homo saviens (man)
	C;Date: 06-	Jul-1982 #seq	nence_re	evision 27-Nov-1985 #te	xt_change 09-Jul-2004
	R;Ohkubo, I	.; Kurachi, K	276; A2	1900; A27699; A31905; A sawa, T.; Shiokawa, H.;	34030 Sasaki, M.
	Biochemistr	/ 23, 5691-56	97, 1984 human ci	4 for alpha-2-thicl p	roteinage inhibitor and its id
	A; Reference	number: A904	90; MUII	D:85122621; PMID:644159	<b>1</b>
	A; Molecule type: mRNA	A; Molecule type: mRNA			
ed,	A;Cross-ref	erences: UNIP	ROT:PO10	042; UNIPARC:UPI000000D	A;Cross-references: UNIPROT:P01042; UNIPARC:UPI000000D92C; GB:K02566; NID:g177889; PIDN R:Takagaki V : Kitamura N : Nakanishi S
	J. Biol. Ch	em. 260, 8601	-8609, 1	1985	
	A;Title: Cit	number: A925	44; MUII	0:85234582; PMID:298929	A;Title: Cloning and sequence analysis of chas for human high molecular weight and low A;Reference number: A92544; MUID:85234582; PMID:2989293
	A; Accession: B25276 A; Molecule type: mRNA	: B25276			
5 1	A; Residues:	A; Residues: 1-427 <tak></tak>	ABC . ITBT	0000000000. CB.W11437.	
× ore	R; Lottspeic	), F.; Keller	mann, J.	.; Henschen, A.; Rauth,	A;cross-rererences: UNIPARC:UPIUUUUUUU92C; GB:MII437; NIU:gI86751; FIUN:AAB59551.1; FIU R;Lottspeich, F.; Kellermann, J.; Henschen, A.; Rauth, G.; Mueller-Esterl, W.
ים ו סיו	in Kining I	/, part A, Gr	eenbaum,	L.M., and Margolius,	in Kinins IV, part A, Greenbaum, L.M., and Margolius, H.S., eds., pp.91-95, Plenum, New
M II D	A;Title: Am A;Reference	A;Title: Amino acid sequen A;Reference number: A27900	00 00	the right chain or hum	an iow moiecular mass kininoge
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hase	Biochem. Bio	ophys. Res. C	ommun.	Biochem. Biophys. Res. Commun. 152, 519-526, 1988	
vine	A; Reference	number: A276	99; MUII	0:88209021; PMID:3365237	7
clone	A; Molecule type: pro	:ype: protein			
ecure rat	A;Residues: A;Cross-ref	A;Residues: 380-389 <min> A;Cross-references: UNIPARC:UPI000002C0F8</min>	ARC:UPIO	000002C0F8	
ff ad	R; Maeda, H.	Matsumura,	Y.; Kato	о, н.	
ecurs ecurs	A;Title: Pu:	em. 263, 1605 rification an	1-16054, d identi	J. BIOL. Chem. 263, 16051-16054, 1988 A; Title: Purification and identification of [hydroxyprolyl(3)] bradykinin in	olyl(3)]bradykinin in ascitic flu¤
recur	A; Reference number: A; Accession: A31905	number: A319 : A31905	05; MUII	D:89034061; PMID:318278	N
A pr	A; Molecule	A; Molecule type: protein	,		
um sa	A;Cross-ref	A; Cross-references: UNIPARC: UPI000002CF4A	ARC: UPIC	00002CF4A	
ecurs	K;Sasaguri, Biochem. Bio	m.; ikeda, m phys. Res. C	ommun. 1	150, 511-516, 1988	
ursor prote	A;Title: Id: A;Reference	ntification number: A340	of [hydz 30; MUII	roxyproline(3)]-lysyl-b D:88106632; PMID:333772	A;Title: Identification of [hydroxyproline(3)]-lysyl-bradykinin released from human plas A;Reference number: A34030; MUID:88106632; PMID:3337729
G 1 6 6 6			****		

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A,Gene: GDB: KNG
A,Gross-references: GDB:125256; OMIM:228960
A,Gross-references: GDB:125256; OMIM:228960
A;Map position: 3q27-3q27
A;Introns: 65/3; 102/3; 131/1; 188/3; 224/3; 253/1; 310/3; 346/3; 375/3; 401/3
C,Superfamily: kininogen; cystatin homology
C;Keywords: alternative splicing; blood coagulation; cysteine proteinase inhibitor; glyc
F;1-18/Domain: signal sequence #status predicted <SIG>
F;1-18/Domain: signal sequence #status predicted <MAT2>
F;19-389,390-427/Product: LMW kininogen (kininogen I) #status predicted <MAT2>
F;19-379/Product: LMW kininogen heavy chain #status predicted <MAT2>
F;19-379/Product: LMW kininogen reference *AFT2>
F;29-379/Domain: cystatin homology <CY1>
F;381-399/Domain: cystatin homology <CY2>
F;381-399/Product: LWW kininogen light chain #status experimental <BDY>
F;381-389/Product: LWW kininogen light chain #status experimental <CH>
F;381-389/Product: LWW kininogen light chain #status experimental <CH>
F;38-407,83-94,107-126,142-145,206-218,229-248,264-267,328-340,351-370/Disulfide bonds:
F;48,169,205,294/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;389-390/Cleavage site: Met-Lys (Kallikrein) #status experimental
F;389/390-390/Cleavage site: Arg-Ser (Kallikrein) #status experimental
F;389-390/Cleavage site: Arg-Ser (Kallikrein) #status experimental
F;401/Binding site: carbohydrate (Thr) (covalent) #status experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Accession: A34030
A;Molecule type: pr
A;Residues: 380-389
C;Species: Homo Bapiens (man)
C;Date: 28-May-1986 #seqmence revision 28-May-1986 #text change 09-Jul-2004
C;Date: 28-May-1986 #seqmence revision 28-May-1986 #text change 09-Jul-2004
C;Accession: A01279; A52276; S32422; A91153; A24871; A27899; A27699; A31905,
R;Ohkubo, I.; Kurachi, K.; Takasawa, T.; Shiokawa, H.; Sasaki, M.
Biochemistry 23, S691-5697, 1984
A;Title: Isolation of a human cDNA for alpha-2-thiol proteinase inhibitor at A;Reference number: A90490; MUID:85122621; PMID:6441591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fed. Proc. 27, 52-57, 1968
A;Title: Structural features of plasma kining and kiningens. A;Reference number: A91455; MUID:90255622; PMID:4952632
A;Contents: annotation; bradykiningens in more of from the contents.
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A;Title: Structural organization of the human kininogen gene and a A;Reference number: A92545; MUID:85234583; PMID:2989294
                                                                                                                                                                                                                                                  kininogen, HMW precursor [validated] - human
N;Alternate names: alpha-2-thiol proteinase inhibitor;
N;Contains: bradykinin (kallidin I); HMW kininogen I; F
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Residues: 380-389 <SAS>
Cross-references: UNIPARC:UPI000002C0F8
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Pred. No. 7.8e-55;
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                                                                                                                                                                                                                                                         preprokininogen;
HMW kininogen II;
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low molecular
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A;Molecule type: protein
A;Molecule type: protein
A;Residues: 381-389 <MAE>
A;Residues: 381-389 <MAE>
A;Cross-references: UNIPARC:UPI000002CF4A
R;Cross-references: UNIPARC:UPI0000002CF4A
R;Sasaguri, M.; Ikeda, M.; Ideishi, M.; Arakawa, K.
R;Sasaguri, M.; Ikeda, M.; Ideishi, M.; Arakawa, K.
Biochem. Biophys. Res. Commun. 150, 511-516, 1988
Biochem. Biophys. Res. Commun. 150, 511-516, 1988
A;Title: Identification of [hydroxyproline(3)]-lysyl-bradykinin released from
A;Title: Identification of [hydroxyproline(3)]-lysyl-bradykinin released from
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A; Molecule type: protein
A; Residues: 379-389,'K',390-407,'Q',409-644 <KEL2>
A; Cross-references: UNIPARC: UPI000017429A
A; Cross-references: UNIPARC: Proud, D.; Walz, D.; Sc
R; Mindrodu, T.; Carretero, O.A.; Proud, D.; Walz, D.; Sc
Biochem. Biophys. Res. Commun. 152, 519-526, 1988
Biochem. Biophys. Res. Commun. 152, 519-526, 1988
A; Title: A new kinin molety in human plasma kininogens.
A; Reference number: A27699; MUID:88209021; PMID:3365237
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A,Note: the bradykinin sequence preceding the light chain sequence R,Kellermann, J.; Lottspeich, F.; Henschen, A.; Mueller-Esterl, W. Eur. J. Biochem. 154, 471-478, 1986
Eur. J. Biochem. 154, 471-478, 1986
A,Title: Completion of the primary structure of human high-molecula A,Reference number: A24871; MUID:86108361; PMID:3484703
A,Accession: A24871
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A;Residues: 1-592,'I',594-644 <TAK>
A;Residues: 1-592,'I',594-644 <TAK>
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M11437;
A;Cross-references: UNIPARC:UPI000013D5AC; GB:M1437;
A;Cross-references: UNIPARC:UPI0000013D5AC; GB:M1437;
A;Cross-re
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J. Biol. Chem. 260, 8601-8609, 1985
A;Title: Cloning and sequence analysis of cDNAs for human high molecular weight and A;Reference number: A92544; MUID:85234582; PMID:2989293
A;Accession: A25276
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A;Residues: 1-389 <OHK>
A;Cross-references: UNIPROT:P01042; UNIPARC:UPI0000174296;
A;Cross-references: UNIPROT:P01042; UNIPARC:UPI0000174296;
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A;Residues: 'Z', 20-380 <KEL1>
A;Cross-references: UNIPARC:UPI0000174299
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R;Lenarcic, B.; Gabrijelcic, D.; Rozman, B.; Drobnic-Kosorok, Biol. Chem. Hoppe-Seyler 369, 257-261, 1988
A;Title: Human cathepsin B and cysteine proteinase inhibitors A;Reference number: S02482; MUID:89076517; PMID:3264507
                                                                                                                                                                                                                                                          A; Molecule type: protein A; Residues: 380-389 < SAS>
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PMID:3484703
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A;Cross-references: GDB:125256; OMIM:228960
A;Map position: 3q27-3q27
A;Introns: 65/3; 102/3; 131/1; 188/3; 224/3; 253/1; 310/3; 346/3; 375/3
A;Introns: 65/3; 102/3; 131/1; 188/3; 224/3; 253/1; 310/3; 346/3; 375/3
C;Superfamily: kininogen; cystatin homology
C;Keywords: alternative splicing; blood cogqulation; cysteine proteinase inhibitor; C;Keywords: alternative splicing; blood cogqulation; cysteine proteinase inhibitor; F;1-18/Domain: sdignal sequence #status experimental <SIG>
F;19-644/Product: HMW kininogen I (prokininogen) #status experimental <MATI>
F;19-379,390-644/Product: HMW kininogen II #status experimental <MATI>
                                                                                                                                                                                                                                                                                                                                                                             C;Comment: The HMW kininogen precursor and the LMW form are produced from the same gene C;Comment: Kininogen is a cysteine proteinase inhibitor, takes part in initiation of the C;Comment: The glycine/histidine/lysine-rich region of HMW kininogen light chain is impor C;Comment: Bradykinin, released from kininogen by kallikrein, is a potent vasodilator, in a potent vasodilator, in a potent vasodilator, in a potent vasodilator, in the kininogen prior to the release of bradykinin.
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A; Residues: 450-452, 'X', 454, 'X', 456 <LIT>
A; Cross-references: UNIPARC: UPI000017429E
R; Straczek, J: Maachi, F: le Nguyen, D:
FEBS Lett. 373, 207-211, 1995
A; Title: Purification from human plasma of
A; Reference number: S68059; MUID: 96033974;
A; Accession: S68059
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A;Residues: 264-359, W., 361-375 <LEN2>
A;Crose-references: UNIPARC:UPI000017429D
R;Little, S.S.; Johnson, D.A.
Biochem. J. 307, 341-346, 1995
A;Title: Human mast cell tryptase isoforms:
A;Reference number: 855239; MUID:95251593; F
A;Accession: 855239
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R;Lenarcic, B.; Krasovec, M.; Ritonja, A.; Olafsson, I.; Tu
FEBS Lett. 280, 211-215, 1991
A;Title: Inactivation of human cystatin C and kininogen by
A;Reference number: S14303; MUID:91192133; PMID:2013314
A;Accession: S14447
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A;Residues: 431-434 <TR>
A;Residues: 431-434 <TR>
A;Cross-references: UNIPARC:UPI000017429F
A;Cross-references: UNIPARC:UPI000017429F
T, Structural N.; Kitagawa, H.; Fukushima, D.; Takagaki, Y.; Miyata, T.; Nakanishi, J. Biol. Chem. 260, 8610-8617, 1985
A;Title: Structural organization of the human kininogen gene and a model for its of A;Reference number: A92545; MUID:85234583; PMID:2989294
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A;Cross-references: UNIPARC:UPI000002CF4A; UNIPARC:UPI0000143284; UNIPARC:UPI000017429B
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A;Experimental source: urine
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A;Cross-references: UNIPARC:UPI000002C0F8
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                                                                                                                                                                                                                                                                                                                   A;Gene: GDB:KNG
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A;Accession: A61495
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     J.V.
27, 52-57, 1968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    252-254,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pro-383 modified to 4-hydroxyproline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a tetrapeptide that potentiates insulin-like PMID:7589467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Becchi, M.; Heulin,
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F;19-379/Domain: HWW kininogen heavy chain #btatus experimental <HCH>
F;19-131/Domain: cystatin homology <CY1>
F;42-253/Domain: cystatin homology <CY2>
F;264-375/Domain: cystatin homology <CY3>
F;264-375/Domain: cystatin homology <CY3>
F;380-389/Product: lysyl-bradykinin (kallidin II) #status experimental <BDY>
F;380-389/Product: bradykinin (kallidin I) #status experimental <CCIP
F;380-389/Product: bradykinin (kallidin I) #status experimental <CCIP
F;380-389/Product: lowmolecular weight growth promoting factor #status experimental
F;41-510/Region: glycine/histidine/lysine-rich 30-residue repeats
F;421-510/Region: glycine/histidine/lysine-rich 30-residue repeats
F;431-434/Product: low molecular weight growth promoting factor #status experimental
F;431-349/Product: low molecular weight growth promoting factor #status experimental
F;48/Binding site: pyrrolidone carboxylic acid (Gln) (in mature form) #status experimental
F;48/Binding site: carbohydrate (Asn) (covalent) #status experimental
F;48/Binding site: carbohydrate (Asn) (covalent) #status experimental
F;389-980/Cleavage site: Met-Lys (kallikrein) #status experimental
F;389-990/Cleavage site: Arg-Ser (kallikrein) #status experimental
F;401,533,542,546,557,571,593,628/Binding site: carbohydrate (Thr) (covalent) #status experimental
F;1-18/Joomain: signal sequence #status predicted <SIG>
F;19-436/Product: LMW kininogen I #status predicted <MAT>
F;19-436/Product: LMW kininogen I heavy chain #status predicted <HCH>
F;19-378/Product: LMW kininogen I heavy chain #status predicted <HCH>
F;19-130/Domain: cystatin homology <CY1>
F;141-252/Domain: cystatin homology <CY2>
F;363-374/Domain: cystatin homology <CY2>
F;363-374/Domain: cystatin homology <CY3>
F;363-388/Product: lysyl-bradykinin (kallidin II) #status predicted <BDY>
F;380-388/Product: LMW kininogen I light chain #status experimental <LCH>
F;380-389/Product: LMW kininogen I light chain #status experimental <LCH>
F;380-389/Product: LMW kininogen I light chain #status experimental <LCH>
F;27-406, 82-93, 106-125, 141-144, 205-217, 228-247, 263-266, 327-339, 350-369/Disulfide
F;378-379/Cleavage site: Mtydroxyproline (Pro) #status predicted
F;382/Modified site: Arg-Ser (kallikrein) #status predicted
F;388-389/Cleavage site: Arg-Ser (kallikrein) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: mRNA
A; Residues: 1-436 < NNA>
A; Residues: 1-436 < NNA>
A; Cross-references: UNIPROT: P01046; UNIPARC: UPI000012DF24; GB: J00010; GB: V00426; NID: g16
C; Comment: The LMW kininogen precursor is produced from the same gene as the HMW form at C; Comment: Kininogen is a cysteine proteinase inhibitor, takes part in initiation of the C; Comment: Bradykinin, released from kininogen by kallikrein, is a potent vasodilator, xyproline residue is present in the kininogen prior to the release of bradykinin.
C; Superfamily: kininogen; cystatin homology
C; Keywords: alternative splicing; blood coagulation; cysteine proteinase inhibitor; glyc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         kininogen, LMW I precursor - bovine
N;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen
N;Contains: bradykinin (kallidin); kininogen I; kininogen II; prokininogen
C;Species: Bos prinigenius taurus (cattle)
C;Date: 14-Nov-1983 #sequence_revision 14-Nov-1983 #text_change 09-Jul-2004
C;Accession: A01283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Nawa, H.; Kitamura, N.; Hirose, T.; Asai, M.; Inayama, Proc. Natl. Acad. Sci. U.S.A. 80, 90-94, 1938
A;Title: Primary structures of bovine liver low molecular A;Reference number: A33984; MUID:83117859; PMID:6572010
A;Accession: A01283
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Pred. No. 1.2e-54;
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kininogen, HMW I precursor - bovine
N;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen
N;Contains: bradykinin (kallidin); kininogen I; kininogen II; prokininogen
C;Species: Bos primigenius taurus (cattle)
C;Bate: 14-Nov-1983 #sequence revision 14-Nov-1983 #text_change 09-Jul-2004
C;Accession: A01281; A91923; A91938; A29559
R;Kitamura, N.; Takagaki, Y.; Furuto, S.; Tanaka, T.; Nawa, H.; Nakanishi, S.
Nature 305, 545-549, 1983
A;Title: A single gene for bovine high molecular weight and low molecular weight kinino, A;Reference number: A93317; MUID:84014106; PMID:6571699
A;Accession: A01281
A;Accession: A01281
A;Cross-references: UNIPROT:P01044; UNIPARC:UPI000012DFIC; GB:V01491; GB:K01757; NID:949
A;Accession: A9323, 1970
U. Biochem. 67, 313-323, 1970
U. Biochem. 67, 313-323, 1970
A;Title: Studies on the structure of bovine kininogen: cleavages of disulfide bonds and A;Reference number: A91923
A;Accession: A91923
A;Accession: A91923
A;Accession: A91923
A;Title: Studies on the primary structure of bovine high-molecular-weight kininogen. Ami A;Reference number: A91938; MUID:75170265; PMID:1169237
A;Molecule type: protein
A;Molecule type: protein
A;Molecule type: protein
A;Molecule type: protein
A;Molecule type: protein
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A; Residues: 458-498 cHAN-
A; Cross-references: UNIPARC: UPI00001742A2
A; Cross-references: UNIPARC: UPI00001742A2
Biol. Chem. 262, 2768-2779, 1987
A; Title: Bovine high molecular weight kininogen. The amino
A; Reference number: A92627; MUID:87137530; PMID:3546295
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                                                                                                                                                                                      Seikagaku 56, 808, 1984
A;Title: Disulfide bonds in bovine
                                                                                                                                                                                                                                                                                                                            A;Title:
                                                                                                                                                                                                                                                                                                                                                    R;Lottspeich, F.; Kellermann, J.; Henschen, A.; Foertsch, Eur. J. Biochem. 152, 307-314, 1985
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;Note: article in Japanese ;Comment: The HMW kininogen precursor is produced from the same gene as the LMW f;Comment: The HMW kininogen precursor is produced from the same gene as the LMW f;Comment: Kininogen is a cysteine proteinase inhibitor, takes part in initiation;Comment: The glycine/histidine/lysine-rich region of HMW kininogen light chain i;Comment: Bradykinin, released from kininogen by kallikrein, is a potent vasodila
                                                                                                                                  Reference number: A94300 ;Contents: annotation; disulfide
                                                                                                                                                                                                                                      Title: The amino acid sequence of the light chain of human high-molecular-mass;Reference number: A91153; MUID:86030270; PMID:4054110;Contents: annotation; bovine cleavage sites; bovine carbohydrate binding sites;Sueyoshi, T.; Miyata, T.; Kato, H.; Iwanaga, S.
                                                                                                                                                                                                                                                                                                                                                                                                      ;Molecule type: protein
;Residues: 'Z',20-123,'I',125-127,'I',129-378
;Cross-references: UNIPARC:UPI00001742A3
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Pred. No. 1.7e-37;
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F;19-130/Domain: Cystatin homology <CY1>
F;141-252/Domain: Cystatin homology <CY2>
F;141-252/Domain: Cystatin homology <CY2>
F;263-374/Domain: Cystatin homology <CY3>
F;263-374/Domain: Cystatin homology <CY3>
F;379-388/Product: lysyl-bradykinin (kallidin I) #status experimental <BDY>
F;389-621/Product: hysyl-bradykinin | #status experimental <CCH>
F;389-621/Product: hysyl-bradykinin (kallidin I) #status experimental <CCH>
F;389-621/Product: hysyl-bradykinin (kallidin I) #status experimental <CCH>
F;389-621/Product: hysyl-bradykinin (kallidin I) #status experimental 

//rroduct: hysyl-bradykinin | #status experimental 

//roduct: hysyl-bradylic acid (Gln) (in mature form) #status experimental F;197/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form) #status experimental F;37,168,169,204/Binding site: carbohydrate (Asn) (covalent) (partial) #status experimental F;378-379/Cleavage site: Met-Lys (kallikrein) #status experimental F;388-389/Cleavage site: Met-Lys (kallikrein) #status experimental F;388-389/Cleavage site: Arg-Ser (kallikrein) #status experimental F;398,406,512/Binding site: carbohydrate (Ser) (covalent) #status experimental F;398,406,520,524,536,548,533,570/Binding site: carbohydrate (Thr) (covalent) #status experimental F;398,400,520,524,536,548,553,570/Binding site: carbohydrate (Thr) (covalent) #status experimental F;398,400,520,524,536,548,553,570/Binding site: carbohydrate (Thr) (covalent) #status experimental F;398,400,520,524,536,548,553,570/Binding site: carbohydrate (Ser) (covalent) #status experimental F;398,400,520,524,536,548,533,570/Binding site: carbohydrate (Ser) (covalent) 
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A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-434 <NAW>
A;Residues: 1-434 <NAW>
A;Cross-references: UNIPROT:P01047; UNIPARC:UPI000012DP28; GB:V00427; GB:J00011; NID:94
C;Comment: The LMW kininogen precursor is produced from the same gene as the HMW form a
C;Comment: Kininogen is a cysteine proteinase inhibitor, takes part in initiation of th
C;Comment: Bradykinin, released from kininogen by kallikrein, is a potent vasodilator,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            kininogen, LMW II precursor - bovine
N;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen
N;Contains: bradykinin (kallidin); kininogen I; kininogen II; prokininogen
C;Species: Bos primigenius taurus (cattle)
C;Date: 14-Nov-1983 #sequence_revision 14-Nov-1983 #text_change 09-Jul-2004
C;Date: 14-Nov-1983 #sequence_revision 14-Nov-1983 #text_change 09-Jul-2004
C;Accession: A01284
R;Nawa, H.; Kitamura, N.; Hirose, T.; Asai, M.; Inayama, S.; Nakanishi, S.
Proc. Natl. Acad. Sci. U.S.A. 80, 90-94, 1983
A;Title: Brinary structures of bovine liver low molecular weight kininogen ]
A;Reference number: A93984; MUID:83117859; PMID:6572010
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                                        F;19-130/Domain: cystatin homology <CYI>
F;141-252/Domain: cystatin homology <CYI>
F;241-252/Domain: cystatin homology <CYIS
F;251-372/Domain: cystatin homology <CYIS
F;377-386/Product: lysyl-bradykinin (kallidin I) #status predicted <KBDY>
F;378-386/Product: bradykinin (kallidin I) #status predicted <BDY>
F;378-386/Product: bradykinin (kallidin I) #status predicted <BDY>
                                                                                                                                                                                                                                                                                                                          C;Superfamily: kininogen; cystatin homology
C;Keywords: alternative splicing; blood coagulation; c;
C;Keywords: alternative splicing; blood coagulation; c;
F;1-18/Domain: signal sequence #status predicted <SIG>
F;19-434/Product: LMW kininogen II #status predicted <I
F;19-377/Product: LMW kininogen I heavy chain #status |
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in, is a potent vasodilator, release of bradykinin.
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A;Molecule type: protein
A;Residues: 456.496 <HA2>
A;Cross-references: UNIPARC:UPI00001742A2
A;Cross-references: UNIPARC:T; Hashimoto, N.; Kato, H.; Hayash
C, Sueyoshi, T.; Miyata, T.; Hashimoto, N.; Kato, H.; Hayash
J, Biol. Chem. 262, 2768-2779, 1987
A;Title: Bovine high molecular weight kininogen. The amino
A;Reference number: A92627; MUID:87137530; PMID:3546295
A;Accession: B29559
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kininogen, HMW II precursor - bovine

k;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen

N;Conteins: bradykinin (kallidin); kininogen I; kininogen II; prokininogen

C;Species: Bos primigenius taurus (cattle)

C;Date: 14-Nov-1983 #sequence revision 14-Nov-1983 #text_change 09-Jul-2004

C;Date: 14-Nov-1983 #sequence revision 14-Nov-1983 #text_change 09-Jul-2004

C;Accession: A01282; A91923; A91941; A91938; B29559

R;Kitamura, N; Takagaki, Y; Furuto, S; Tanaka, T; Nawa, H; Nakanishi, Nature 305, 545-549, 1983
A;Molecule type: protein
A;Residues: 'Z',20-104,'E',106-256,'XX',257-376
A;Cross-references: UNIPARC:UPI00001742A5
                                                                                                                                                                                                                                                                                                                              A,Title: Studies on the primary structure of bovine high-molecular-weight kininogen. A,Reference number: A91938; MUID:75170265; PMID:1169237 A,Accession: A91938
                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: UNIPARC:UPI00001742A4
A;Note: 398-Pro, 401-Val, and 455-Lys were also found
R;Han, Y.N., Komiya, M.; Iwanaga, S.; Suzuki, T.
J. Biochem. 77, 55-68, 1975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Han, Y.N.; Kato, H.; Iwanaga, S.; Suzuki, T.
J. Biochem. 79, 1201-1222, 1976
A;Title: Primary structure of bovine plasma high-molecular-weight kininogen.
A;Reference number: A91941; MUID:76260155; PMID:956151
A;Accession: A91941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: mRNA
A;Residues: 1-619 <KIT>
A;Residues: 1-619 <KIT>
A;Cross-references: UNIPROT:P01045; UNIPARC:UPI000012DF1F;
A;Cross-references: UNIPROT:P01045; UNIPARC:UPI000012DF1F;
A;Kato, H.; Nagasawa, S.; Suzuki, T.
Biochem. 67, 313-323, 1970
J. Biochem. 67, 313-323, 1970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Reference number: A93317; MUID:84014106; PMID:6571699 A;Accession: A01282
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A,Residues: 376-391 <KAT>
A,Cross-references: UNIPARC:UPI00001742A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A,Title: Studies on the structure of bovine A,Reference number: A91923; MUID:70180420; PA,Accession: A91923
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Matches 84
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         252 GEDFL--PPMVCVGCPKPIPVDSPDLEEALNHSIAKLNAEHDGTFYFKIDTVKKATVQVV
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Pred. No. 6.2e-35;
4; Mismatches 22; Indels
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PMID:4986212
                                                                                                                                                                                                          H.; Hayashida,
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F;141-252/Domain: Cystatin homology <CY2>
F;261-372/Domain: Cystatin homology <CY3>
F;261-372/Domain: Cystatin homology <CY3>
F;377-386/Product: lysyl-bradykinin (kallidin II) #status experimental <RBDY>
F;378-386/Product: bradykinin (kallidin I) #status experimental <LCH>
F;387-589/Product: bradykinin (kallidin I) #status experimental F;19/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form) #status experiment
F;27-589,82-93,106-125,141-144,205-217,228-247,261-264,325-337,348-367/Disulfide bonds:
F;377-168,169,204,280/Binding site: carbohydrate (Asn) (covalent) #status experimental
F;376/Binding site: carbohydrate (Asn) (covalent) #status experimental
F;376/Binding site: carbohydrate (Asn) (covalent) #status experimental
F;376/Binding site: 4-hydroxyproline (Pro) #status experimental
F;386-387/Cleavage site: Arg-Ser (kallikrein) #status experimental
F;397,398,518,522,534,546,551,588/Binding site: carbohydrate (Ser) (covalent) #status experimental
F;397,398,518,522,534,546,551,588/Binding site: carbohydrate (Thr) (covalent) #status experimental
F;397/Cleavage site: Arg-Thr (kallikrein) #status experimental
                            A;Reference number: A92496;
A;Accession: A28055
A;Molecule type: mRNA
                                                                                                                                                                                                                                                                 K-kininogen, LMW I precursor - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 20-Jun-1989 #sequence_revision 20-Jun-1989 #text_change
C;Accession: A28055
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C;Superfamily: kininogen; cystatin homology
C;Keywords: alternative splicing; blood coagulation; cysteine proteinase inhibitor; dup
                                                                                                                                                                                                           R;Furuto-Kato, S.; Matsumoto, A.; Kitamura, J. Biol. Chem. 260, 12054-12059, 1985
                                                                                                                                                                                                                                                                                                                                                                                A28055
K-kininogen,
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F;19-619/Product: HMW kininogen II #status predicted <MAT-
F;19-376/Product: HMW kininogen II heavy chain #status exp-
F;19-130/Domain: cystatin homology <CY1-
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A;Title: Disulfide bonds in bovine HMW kininogens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eur. J. Blochem. 152, 307-314, 1985
A;Title: The amino acid sequence of the light chain of human high-molecular-mass
A;Reference number: A91153; MUID:86030270; PMID:4054110
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1-433 <FUR>
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68.9%;
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No. 9.1e-35;
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references: UNIPROT: P08934; UNIPARC: UPI0000177A66

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N;Contains bradykinin
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 08-Mar-1989 #sequence_revision 08-Mar-1989 #text_change 09-Jul-2004
C;Accession: A25486
R;Kitagawa, H.; Kitamura, N.; Hayashida, H.; Miyata, T.; Nakanishi, S.
J;Biol. Chem. 262, 2190-2198, 1987
J,Title: Differing expression patterns and evolution of the rat kininogen gene family.
A;Title: Differing expression patterns and evolution of the rat kininogen gene family.
A;Reference number: A92625; MUID:87137443; PMID:3029068
A;Accession: A25486
A;Molecule type: mRNA
A;Residues: 1-639 <KIT>
A;Cross-references: UNIFROT:P08934; UNIFARC:UPI000012DF1B
A;Note: the authors translated the codon CAA for residue 347 as Asn
C;Superfamily: kininogen; cystatin homology
C;Keywords: alternative splicing
F;1-18/Domain: signal sequence #status predicted <MAT>
F;19-639/Product: kininogen, HMW I #status predicted <MAT>
F;19-131/Domain: cystatin homology <CV2>
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major acute phase alpha-1 protein (version C;Species: Rattus norvegicus (Norway rat) C;Date: 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 19-May-1989 #sequence_revision 
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A23897
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   Species: Rattus norvegicus (Norway rat)

Date: 19-May-1989 #sequence_revision 19-May-1989 #text_change 09-Jul-2004;

Accession: A23897; B23897
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264-375/Domain: cystatin homology <CY3>
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; Pred. No. 1.2e-32;
14; Mismatches 29
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Pred. No. 1.8e-32;
4; Mismatches 29;
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 20-Jun-1989 #sequence revision 20-Jun-1989 #text_change 09-Jul-2004
C;Accession: B28055; E25486; B28526; C28526
R;Furuto-Kato, S.; Matsumoto, A.; Kitamura, N.; Nakanishi, S.
J. Biol. Chem. 260, 12054-12059, 1985
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Note: the authors translated the codon C;Superfamily: kininogen; cystatin homolc F;19-130/Domain: cystatin homology <CY1: F;141-252/Domain: cystatin homology <CY2:
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A;Residues: 5-430 <AND2>
A;Cross-references: UNIPARC:UPI0000170A0C;
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A;Accession: B23897
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A;Title: The relationship between rat major acute phase A;Reference number: A23897; MUID:86008266; PMID:2413019
A;Accession: A23897
A;Molecule type: protein
A;Residues: 1-14 <AND1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPROT:P08932; UNIPARC:UPI000012DF30
R;Kitagawa, H.; Kitamura, N.; Hayashida, H.; Miyata, T.; N
J. Biol. Chem. 262, 2190-2198, 1987
A;Title: Differing expression patterns and evolution of th
A;Reference number: A92625; MUID:87137443; PMID:3029068
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                                                                                               A; Molecule type: protein
A; Residues: 'E', 20-48; 376-388, 'R', 390-419, 'ER', 422-430
A; Cross-references: UNIPARC: UPI0000177A67
                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPARC:UPI0000000380
R;Enjyoji, K.; Kato, H.; Hayashi, I.; Oh-ishi, S.; Iwanaga, S.
J. Biol. Chem. 263, 973-988
A;Title: Purification and Characterization of rat T-kininogens
A;Reference number: A92729; MUID:88087226; PMID:3121623
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                                                                                                                                                                                                 A;Molecule type: protein
A;Residues: 'E',20-25,'MD',28-48,376-430 <ENJ>
A;Cross-references: UNIPARC:UPI0000177A67
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A; Residues: 1-430 < FUR>
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A; Residues: 375-430 < KIT>
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Best Local
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                            #status predicted <SIG>
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Pred. No. 4.7e-31;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    evolution of the rat kininogen PMID:3029068
     experimental <MAT>
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A;Reference number: A0128
A;Accession: A01285
Å;Molecule type: mRNA
A;Residues: 1-423 <COL>
A;Cross-references: UNIPF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Cole, T.; Inglis, A.S.; Roxburgh, C.M.; Howlett, G.J.; Schreiber, FBBS Lett. 182, 57-61, 1985
"A;Title: Major acute phase alphal-protein of the rat is homologous A;Reference number: A01285; MUID:85127561; PMID:2578992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        r.
C;Superfamily: kininogen; cystatin homology
C;Keywords: bradykinin; cysteine proteinase
P;1-11/Domain: signal sequence (fragment) #c
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F;141-252/Domain: cystatin homology <CY2>
F;141-252/Domain: cystatin homology <CY2>
F;263-374/Domain: cystatin homology <CY2>
F;19/Modified site: pytrolidone carboxylic acid (Gln) (in mature form) #status experimen F;19/Modified site: pytrolidone carboxylic acid (Gln) (covalent) #status predicted F;82,126,168,204,326/Binding site: carbohydrate (Asn) (covalent) #status predicted F;83-94,107-125,141-144,205-217,228-247,263-266,327-339,350-369/Disulfide bonds: #status
T-kininogen
N,Alternate
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
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136-367/Domain: cystatin homology <CY3>
1371-379/Product: bradykinin #status predicted <BDY>
12/Modified site: pyrrolidone carboxylic acid (Gln)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cross-references: UNIPARC:UPI00001742A6 Comment: This plasma glycoprotein inhibits cysteine proteinases. During acute
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Species: Rattus norvegicus (Norway rat)
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Best Local
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I precursor names: 73K p
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                                                                                                                                                                                                                                                                           AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 122
                                                                                                                                                                                                                                                                                                                       GDDLFELLPKNCRGCPREIPVDSPELKEALGHSIARLNAQHNHIFYFKIDTVKKATSQVV
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                                                                                                                                                                                        LGM 125
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  protein;
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                       rat
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  kininogen
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A;Experimental source: strain Sprague-Dawley R;Kageyama, R.; Kitamura, N.; Ohkubo, H.; Nakanishi, S. J. Biol. Chem. 262, 2345-2351, 1987
A;Title: Differing utilization of homologous transcription A;Reference number: A25488; MUID:87137465; PMID:3818598
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A;Residues: 330-420,'R', 422-429,'P' <KAN>
A;Residues: 330-420,'R', 422-429,'P' <KAN>
A;Cross-references: UNIPARC:UPI00001742A9
R;Anderson, K.P.; Croyle, M.L.; Lingrel, J.B.
Gene 81, 119-128, 1989
Gene 81, 119-128, 1989
A;Title: Primary structure of a gene encoding rat T-kininogen.
A;Reference number: JQ0027; MUID:90034172; PMID:2806908
A;Accession: JQ0027
A,MC1001175.
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A;Molecule type: mRNA
A;Residues: 1-430 <FUR>
A;Residues: 1-430 <FUR>
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references: UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references:UPI0000163B79;
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 17-Mar-1987 #sequence revision 17-Mar-1987 #text_change 31-Dec-2004
C;Date: 10-Mar-1987 #sequence revision 17-Mar-1987 #text_change 31-Dec-2004
C;Accession: A01286; D25486; A28526; PL0193; JQ0007; B25488; A28525; S68036
R;Furuto-Kato, S.; Matsumoto, A.; Kitamura, N.; Nakanishi, S.
J. Biol. Chem. 260, 12054-12059, 1985
                                                                                                                                                                                                                                                                                                                    A;Molecule type: protein
A;Residues: 376.430 cENDE
A;Cross-references: UNIPARC:UPI00001742A8
A;Cross-references: UNIPARC:UPI00001742A8
A;Sierra, F.; Walter, R.; Vautravers, P.; Guigoz, Y.
R;Sierra, F.; Walter, R.; Vautravers, 1995, Guigoz, Y.
A;Ch. Biochem. Biophys. 322, 333-338, 1995
A;Tille: Identification of several isoforms of T-kininogen
A;Reference number: S68034; MUID:96032652; PMID:7574705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPARC:UPI000000041B; GB:M14356; NID:g205090; PIDN:AAA41492. R;Enjyoji, K.; Kato, H.; Hayashi, I.; Oh-ishi, S.; Iwanaga, S. J. Biol. Chem. 263, 965-972, 1986 A;Title: Purification and characterization of two kinds of low molecular weight k A;Reference number: A28525; MUID:88087225; PMID:3335530
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A;Title: Identification of a protein increasing in serum of Nagase analbuminemic rats A;Reference number: PL0193; MUID:90216390; PMID:2108948
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A; Residues: 375-430 < KIT>
A; Cross-references: UNIPARC: UPI00000E7F29
A; Cross-references: UNIPARC: UPI00000E7F29
R; Enjyoji, K.; Kato, H.; Hayashi, I.; Oh-ishi, S.; Iwanaga, S.
J. Biol. Chem. 263, 973-979, 1988
J. Biol. Chem. 263, 973-979, 1988
A; Title: Pwrification and Characterization of rat T-kininogens
A; Reference number: A92729; MUID:88087226; PMID:3121623
A;Cross-references: UNIPARC:UPI00001742AB
A;Experimental source: clone pSG17
C;Comment: At least three types of LMW ki.
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A;Residues: 1-60,'E',62-113,'R',115-165,'F',167-178,'TKI',182-211,'F',213-256,'S',258-3:
A;Cross-references: UNIPARC:UPI00001742AA
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J. Biol. Chem. 262, 2190-2198, 1987
A;Title: Differing expression patterns and
A;Reference number: A92625; MUID:87137443;
                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 340-430 <SIE>
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A; Residues: 1-48 < KAG>
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R;Kanda, S.; Sugiyama, K.; Takahashi, M.; Shumiya, S.; Tomino, S
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A; Residues: 'E', 20-48; 376-430 < ENJ>
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A;Cross-references: UNIPARC:UPI0000173259
C;Superfamily: cystatin; cystatin homology
C;Keywords: colostrum; cysteine proteinase
F;2-112/Domain: cystatin homology <CYS-
F;48-52/Region: inhibitory #status predicte
F;66-76,90-110/Disulfide bonds: #status pre
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F;263-374/Domain: cystatin homology <CY3>
F;378-386/Product: bradykinin #status predicted <BDY>
F;378-386/Product: bradykinin #status oredicted <BDY>
F;19/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form) #status experimen F;82,126,168,204,326/Binding site: carbohydrate (Asn) (covalent) #status predicted F;82,126,168,204,326/Binding site: carbohydrate (Asn) (covalent) #status predicted F;83-94,107-125,141-144,205-217,228-247,263-266,327-339,350-369/Disulfide bonds: #status
                                        RESULT
S68035
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C/Species: Bos primigenius taurus (cattle)
C/Date: 28-Feb-1986 #sequence_revision 28-Feb-1986 #text_change 06-Dec-1996
C/Accession: A01271
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T-kininogen (clone pSG17) - rat (frac
C;Species: Rattus norvegicus (Norway
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R; Hirado, M.; Tsunasawa,
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Best Local S
Matches 76
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Best Local :
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41-45, 1985
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ilarity 30.6%;
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2; Mismatches 31
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Pred. No. 2.7e-30;
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C;Date: 12-Feb-1998 #sequence_revision 20-Feb-1998 #text_change 31-Dec-2004 C;Accession: S68035  
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A;Title: Identification of several isoforms of T-kininogen expressed in the A;Reference number: S68034; MUID:96032652; PMID:7574705  
A;Accession: S68035  
A;Molecule type: mRNA A;Residues: 1-91 <SIE> A;Cross-references: UNIPROT:P70517; UNIPROT:Q63581; UNIPARC:UPI0000177A60  
A;Cross-references: UNIPROT:P70517; UNIPROT:Q63581; UNIPARC:UPI0000177A60
                                                                                                                                                                                                                                                                                                                                                                                         T-kininogen (clone pSG22) - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
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A; Title: Identification of several isoforms of T-kininogen
A; Reference number: S68034; MUID:96032652; PMID:7574705
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Raha Y.J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Raha Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Raha Mhiting M., Madan A., Shevchenko Y., Bouffard G.G.,

Raha Mhiting M., Touchman J.W., Green E.D., Dickson M.C.,

Raha Mitterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

Raha S.Chnerch A., Schein J.E., Jones S.J.M., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human

Tand mouse cDNA sequences.";
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21-JUL-1986 (Rel. :

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Submitted (OCT-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; BC060039; AAH60039.1; -; mRNA.
GO; GO:0004869; F:cysteine protease inhibitor activity;
InterPro; IPR000010; Prot inh_cystat.
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Eukaryota; Metazoa;
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Kato H., Mateumura Y., Maeda H.;
"Isolation and identification of hydroxyproline arm"
TEBS Lett. 232:252-254/77
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Toth E.J., Nickerson D.A.;
"SeattleSNPs. NHLBI HL66682 program for genomic applications,
"SeattleSNPs. WA (URL: http://pga.gs.washington.edu).";
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MEDLINE=85122621, Pu
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Kininogen-1 precursor (Alpha-2-thiol proteinase
Kininogen-1 heavy chain; Bradykinin (Kallidin I)
(Kallidin II); Kininogen-1 light chain; Low mole
promoting factor].
Name=KNG1; Synonyms=BDK, KNG;
                                                                                                                                                                                                       MEDLINE-86108361; PubMed=3484703;
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"Completion of the primary structure of human hig
kininogen. The amino acid sequence of the entire
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Mueller-Esterl
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MEDLINE=90255622;
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            "Purification from human plasma insulin-like growth factor-I act
                                                                                                                                                                                                                                                                                       PROTEIN SEQUENCE OF 19-380, CARBOHYDRATE-LINKAGE ASN-205, AND ABSENCE OF CARBOHYDRATE AT ASN-48.
                                                                                                                                                                                                                                                                                                                                              Pierce J.V.;
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                                        Nabet P.,
                                                   Straczek J.,
                                                                  MEDLINE=96033974;
insulin-like growth factor-I
FEBS Lett. 373:207-211(1995)
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                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid sequence of the light
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J. Biochem.
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                                         Belleville F.;
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                                                     Maachi F., Le Nguyen
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                                                   DF 431-434, AND MASS SPECTROMETRY.
PubMed=7589467; DOI=10.1016/0014-5793
hi F., Le Nguyen D., Becchi M., Heulin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    [GENOMIC DNA]
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              activity
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              of a tetrapeptide that potentiates tivity in chick embryo cartilage.";
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C -!- FUNCTION: (1) Kininogens are inhibitors of thiol proteases; (2)

C -!- FUNCTION: (1) Kininogens are inhibitors of thiol proteases; (2)

C -!- FUNCTION: (1) Kininogens are inhibitors of thiol proteases; (2)

C --- FUNCTION: (1) Kininogen inhibitors of thiologens are too congulation by

C --- FUNCTION: (2) HWW-kininogen inhibits the thrombin-and plasmin-

C --- Function of thrombocytes; (4) the active peptide

C --- Function of thrombocytes; (4) the active peptide

C --- Function of thrombocytes; (4) the active peptide

C --- Function of thrombocytes; (4) natriures and

C --- Function of thrombocytes; (4) it is a

C --- Function of inflammation of hypotension, (4C) natriures and

C --- Function of inflammation and causes (4E1) increase in vascular

C --- Function of inflammation of nociceptors (4E3) release of

C --- Function of inflammation (2) prostaglandins), (4F) it has

C --- Function of inflammation (2) prostaglandins (5)

C --- Function of inhibits the aggregation of thrombocytes; (6) LMW-

LMM-kininogen is in contrast to HWM-kininogen not involved in blood
     EMBL; K02566;
EMBL; M11437;
EMBL; M11438;
                                                                                                                                                                    between
the Euro
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MEDLINE-22660472; PubMed-12754519; DOI=10.1038/nbt827;
Zhang H., Li X.-J., Martin D.B., Aebersold R.;
"Identification and quantification of N-linked glycoprophydrazide chemistry, stable isotope labeling and mass pages. Biotechnol. 21:660-666(2003).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AMINO-ACID COMPOSITION OF 381-389, AND HYDROXYLATION SITE PRO-3 MEDLINE=89034061; PubMed=3182782; Maeda H., Matsumura Y., Kato H.; Merication and identification of [hydroxyprolyl3]bradykinin ascitic fluid from a patient with gastric cancer."; J. Biol. Chem. 263:16051-16054(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sueyoshi T., Miyata T., Kato H., Iwanaga : "Disulfide bonds in bovine HMW kininogens Seikagaku 56:808-808(1984).
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Kitamura N., Kitagawa H., Fukushima
Nakanishi S.;
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J. Biol. Chem.
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"Structural organization of the human
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SIMILARITY: Contains
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SUBCELLULAR LOCATION: Secreted; extracellular.
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                                                                                                                             Swiss-Prot entry is copyright. It is produced through a c
sen the Swiss Institute of Bioinformatics and the EMBL
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Hydroxylation
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HGNC; HGNC:6383; KNG1.
Reactome; P01042; --
MIM; 228960; -
G0:0008201; F:hepsi
G0; G0:0008201; F:rece
G0; G0:0008270; F:rzin
G0; G0:0008270; F:zin
G0; G0:0006954; P:infil
G0; G0:0030146; P:infil
G0; G0:0030147; P:negi
G0; G0:0030147; P:negi
G0; G0:0030185; P:negi
G0; G0:0007162; P:negi
G0; G0:0003165; P:negi
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EMBL; M11528; AAB59550.1; JU

EMBL; M11437; AAB59551.1; JU

EMBL; M11438; AAB59551.1; JU

EMBL; M11521; AAB59551.1; JU

EMBL; M11523; AAB59551.1; JU

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EMBL; M11528; AAB59551.1; JU

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EMBL; M11529; AAB59551.1; JU

EMBL; M1529; AAB69551.1; JU

EMBL; M1529; AAB69551.1; JU

EMBL; M1529; KGHUH.
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GO; GO:000201; F:heparin binding; NAS.
GO; GO:0008201; F:receptor binding; IPI.
GO; GO:0008270; F:zinc ion binding; NAS.
GO; GO:0008270; F:zinc ion binding; NAS.
GO; GO:0008270; F:zinc ion binding; NAS.
GO; GO:0030146; P:diuresis; TAS.
GO; GO:0030147; P:natriuresis; TAS.
GO; GO:0030147; P:negative regulation of blood coagulation; GO; GO:000195; P:negative regulation of cell adhesion; IDI
GO; GO:0043065; P:positive regulation of apoptosis; NAS.
GO; GO:0006939; P:megative regulation of apoptosis; NAS.
GO; GO:0006939; P:megative regulation of apoptosis; NAS.
                                                                                                                                                                                                                                                                                                                                              Vasoactive; Vasodilator.
SIGNAL 1 18
CHAIN 19 644
                                                                                                                                                                                                                                                                                                                                                                                                    Alternative splicing; Blood coagulation; Bradykinin; Direct protein sequencing; Glycoprotein; Hydroxylation; Inflammatory response; Plasma; Polymorphism; Protease i
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InterPro; IPR000010; Prot inh c
Pfam: PF00031; Cystatin; 3.
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M11522;
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M11524;
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M11526;
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SM00043; CY; 3.
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                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                              AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
                                                                                                                                                                                                                                          GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                                                                                                  E
                                                                                                                                                                AGKKYFIDFVARETT
                                                                                                                                                                                                                         GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                         carboxylic acid;
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AAB59550.
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KNG1\_BOVIN ST/ P01044; P01046; 21-JUL-1986 (Rel.

STANDARD;

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BOVIN

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RP PROTEIN SEQUENCE OF 458-498.

RR MEDLINE-75170265; PubMed=1169237;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RI Sinchem. 77:55-68(1975).

CC IP PROCTION: (1) Kininogens are inhibitors of thiol proteases; (2)
CC IP PROCTION: (1) Kininogens are inhibitors of thiol proteases; (2)
CC HAWN-kininogen plays an important role in blood coasgulation by helping to position optimally prekallikrein and factor XI next to factor XII; (3) HMW-kininogen inhibits the thrombin-and plasmin-cc factor XII; (3) HMW-kininogen inhibits the thrombin-and plasmin-cc induced aggregation of thrombocytes; (4) the active peptide bradykinin that is released from HMW-kininogen shows a variety of contraction, (4B) induction of hypotension, (4C) natriuresis and diuresis, (4D) decrease in blood glucose level, (4E) it is a cc mediator of inflammation and causes (EE) increase in vascular permeability, (4E2) stimulation of nociceptors (4E3) release of cother mediators of inflammation (e.g. prostagalandins), (4F) it has a cardioprotective effect (directly via bradykinin action, indirectly via endothelium-derived relaxing factor action); (5) LMW-kininogen is in contrast to HMW-kininogen not involved in blood contraction is in contrast to HMW-kininogen not involved in blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kato H., Nagasawa S., Suzuki T.;
"Studies on the structure of bovine kininogen: cleavages obonds and of methionyl bonds in kininogen-II.";
J. Biochem. 67:313-323(1970).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sueyoshi T., Miyata T., Hashimoto N., Kato H., Miyata T., Iwanaga S.; "Bovine high molecular weight kininogen. The ampositions of carbohydrate chains and disulfide chain portion.":
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13-SEP-2005 (Rel. 48, Last annotation update)
Kininogen-1 precursor (Kininogen I) (Thiol proteinase inhibitor)
[Contains: Kininogen-1 heavy chain; Bradykinin (Kallidin I); Lys
bradykinin (Kallidin II); Kininogen-1 light chain].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-83117859; PubMed-6572010;
Nawa H., Kitamura N., Hirose T., Asai M., In
"Primary structures of bovine liver low mole
precursors and their two mRNAs.";
Proc. Natl. Acad. Sci. U.S.A. 80:90-94(1983)
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Kitamura N., Takagaki Y., Furuto S., Tana
"A single gene for bovine high molecular
weight kininogens.";
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MEDLINE=87137530; PubMed=3546295;
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- ALTERNATIVE PRODUCT
EVENT=Alternative E
Name=HMW;
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IsoId=P01044-1;
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a; Eutheria; Laura
Bovidae; Bovinae;
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                                                                                      LOCATION:
PRODUCTS:
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Laurasiatheria;
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Sequence=Displayed
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heria; Cetartiodactyla;
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cular weight
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v molecular
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     S., Nakanishi S.; weight kininogen
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ulrect protein sequencing; Blood coagulation; Bradykinin;
plasma; Protease inhibitor; Pyrrolidone carboxylic acid; Repeat:
Signal; Thiol protease inhibitor; Vasoactive; Vasodilaria Signal; Thiol protease inhibitor; Vasoactive; Vasodilaria SignAL 1 18 "robable."
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CARBOHYD
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EMBL; V00426; CAA23709.1;
PIR; A01281; KGBOH1.
PIR; A01283; KGBOL1.
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   CONFLICT
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                                     VARSPLIC
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PTM: Bradykinin is released from kininogen by plas SIMILARITY: Contains 3 cystatin-like domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pro; IPR002395; Kininogen.
Pro; IPR000010; Prot_inh_cystat.
Pr00031; Cystatin; 3.
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       ¥.
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Cystatin-like 2.
Cystatin-like 3.
Cleavage (by kallikrein)
Cleavage (by kallikrein)
Cleavage (by kallikrein)
Cleavage (by kallikrein)
Pyrrolidone carboxylic a
N-linked (GlCNAC. .)
N-linked (GlCNAC. .)
N-linked (GlCNAC. .)
N-linked (GlCNAC. .)
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Kininogen-1 light
                                    VSLPHSAMSPVQDEERDSGKEQGPTHGHGWDHGKQI
HVKSCEYKGRPQEAGAEPAPQGEVSLPAESPQLAR
180fOrm LMW).
/FTId=VSP 013562.
Missing (In 180form LMW).
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                          FTId=VSP
        D16850BEFE3C55CD CRC64
                  VSP_013563.
(in_Ref. 2
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(by kallikrein).
(by kallikrein).
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6 outstation -
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Best Local
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Mammalia, Eutheria, L
Pecora, Bovidae, Bovi
NCBI TaxID-9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Miyata T., Wanaga S.;
"Bovine high molecular weight kininogen.
positions of carbohydrate chains and dischain portion.";
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P0.1045; P0.1047;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
Kininogen-2 precursor (Kininogen II) (Thiol proteinase inhibitor)
[Contains: Kininogen-2 heavy chain; Bradykinin (Kallidin I); Lysy
bradykinin (Kallidin II); Kininogen-2 light chain].
                      PROTEIN SEQUENCE OF 387-455.

MEDLINE-76260155; PubMed-956151;

Han Y.N., Kato H., Iwanaga S., Suzuki T.,

"Primary structure of bovine plasma high-molecular-weight kininogen.

"Pame anino acid sequence of a glycopeptide portion (fragment 1)
following the C-cerminus or the bradykinin moiety.";

J. Biochem. 79:1201-1222(1976).
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Nature
[2]
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Proc. Natl. Acad. Sci. U.S.A. 80:90-94(1983).
                                                                                                                                                                                                                                                                                              "Studies on
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MEDLINE=70180420;
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MEDLINE=83117859; PubMed=6572010;
NAWA H., Kitamura N., Hirose T., Asai M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE [MRNA] (ISOFORM HMW).
MEDLINE=84014106; PubMed=6571699;
Kitamura N., Takagaki Y., Furuto S., Tanaka T., Nawa H., Nakanishi S.;
"A single gene for bovine high molecular weight and low molecular
"A single gene for bovine high molecular weight and low molecular
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Nagasawa S., Suzuki T.;
Nagasawa S., Suzuki T.;
on the structure of bovine kininogen: of the structure of bovine kininogen-II.";
d of methionyl bonds in kininogen-II.";
nem. 67:313-323(1970).
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Pred. No. 4.6e-36;
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EMBL; V01492; CAA24736.1; EMBL; V00427; CAA23710.1; PIR; A01282; KGBOH2. PIR; A01284; KGBOL2. HSSP; P01038; 1A90.
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                                                                                                                                urrect protein sequencing; Glycoprotein; Hydroxylation; Inflammatory response; Plasma; Protease inhibitor; Pyrrolidone carboxylic acid; Repeat; Signal; Thiol protestive; Vasoactive; Vasodilator.
                                                                                                                                                                                                                                InterPro; IPR002395; Kinino; InterPro; IPR000010; Prot ii Pfam; PF00031; Cystatin; 3. PRINTS; PR00334; KININOGEN. SMART; SM00043; CY; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-75170265; PubMed-1169237;
MEDLINE-75170265; PubMed-1169237;
Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
"Studies on the primary structure of bovine h
kiningen. Amino acid sequence of a fragment
peptide') released by plasma Kallikrein.";
J. Biochem. 77:55-68(1975).
                                                                                                                               Vasoactive;
SIGNAL
                                                                                                                                                                                   PROSITE; PS00287; CYSTATIN; 2.
Alternative splicing; Blood coagulation;
Direct protein sequencing; Glycoprotein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          helping to position optimally prekallikrein and factor XI next to factor XII; (3) HAW-kininogen inhibits the thrombin-and plasmin-induced aggregation of thrombocytes; (4) the active peptide bradykinin that is released from HAW-kininogen shows a variety of physiological effects: (4A) influence in smooth muscle contraction, (4B) induction of hypotension, (4C) natriuresis and diuresis, (4D) decrease in blood glucose level, (4E) it is a mediator of inflammation and causes (4E) increase in vascular permeability, (4E2) stimulation of nociceptors (4E3) release of other mediators of inflammation (e.g. prostaglandins), (4F) it has a cardioprotective effect (directly via bradykinin action, indirectly via endothelium-derived relaxing factor action); (5) LMW-kininogen is in contrast to HMW-kininogen not involved in blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Iwanaga S., Kato H., (In) Tschesche H., Ho Biological functions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IsoId=P01045-2; Sequence=VSP_013564, VSP_013565; TISSUE SPECIFICITY: Plasma.
PTM: Bradykinin is released from kininogen by plasma kallikrein.
SIMILARITY: Contains 3 cystatin-like domains.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Event=Alternative Name=HMW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FUNCTION: (1) Kininogens are inhibitors of thiol proteases; HMW-kininogen plays an important role in blood coagulation)
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H., Sugo T., Ikari
, Hotzer H. (eds.);
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tin; 3.
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Lysyl-bradykinin.
Bradykinin.
Kininogen-2 light
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Cystatin-like 2.
Cystatin-like 2.
Cystatin-like 3.
Not glycosylated.
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                                                                                                 Kininogen-2
Kininogen-2
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27, Last sequence update)
29, Last sequence update)
29, Last annotation update)
(High molecular weight kininogen I isoform
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Cleavage (by kallikrein).
Cleavage (by kallikrein).
Pyrrolidone carboxylic acid.
4-hydroxyproline (By similarity).
N-linked (GlCNAC...); partial.
O-linked (GalNAC...); partial.
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Missing (In isoform )
/FTId=VSP_013565.
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14; Mismatches
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STANDARD; PRT; 661 AA.

AC 008677; 008676; Q91XK5;

DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 13-SEP-2005 (Rel. 48, Last annotation update)
DT 13-SEP-2005 (Rel. 48, Last annotation update)
DE Kininogen-1 light chain].
DE Kininogen-1 light chain].
DE Kininogen-1 Synonyms=Kng;
SM Name=Kng1; Synonyms=Kng;
SM Namesia; Evaperia; Eva
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Biol. Chem.
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GO; GO:0005515; C:extracellular space; GO; GO:0005515; C:extracellular space;
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Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
NCBI_TaxID=10090;
NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA] (ISOFORM LMW).
STRALN=CS7BL/60; TISSUE=Placenta;
MEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kon
Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=C57BL;
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                                                                                                                                                                                                                                                           STRAIN=C57BL/6 X CBA; TISSUE=Liver;
MEDLINE=97342556; PubMed=9199253; DOI=10.1016/S0167-4781(97)00018-3;
Takano M., Kondo J., Yayama K., Otani M., Sano K., Okamoto H.;
Takano M., Kondo J., Yayama K., Otani M., Sano K., Okamoto H.;
"Molecular cloning of cDNAma for mouse low-molecular-weight and high-
molecular weight prekininogens.";
Biochim. Biophys. Acta 1352:222-230(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE [MRNA] (ISOFORMS STRAIN=C57BL/6 X CBA; TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    122
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to the EMBL/GenBank/DDBJ
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Pred. No. 3.8e
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RA Strausberg R.D., Collins F.S., Wagner L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K.J., Farmer A.A., Rubin G.M., Hong L.,
RA Stepleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., McWann B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Raha S.S., McWann P.J., McKernan K.J., Malek J.A., Gunarathe P.H.,
RA Hiton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Willalon D.K., Munny D.M., Sodergren E.J., Lu X., Gunarathe P.H.,
RA Willalon D.K., Munny D.M., Sodergren E.J., Lu X., Gunarathe P.H.,
RA Hiton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Willalon B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Willalon B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Hopkins R.M., Touchman J.M., Schmutz J., Myers R.M.,
RA Hottried M., Schein J.E., Jones S.J.M., Marra M.A.,
RA Hottried A.S., Schein J.E., Jones S.J.M., Marra M.A.,
RA Hottried A., Schein J.E., Jones S.J.M., Marra M.A.,
RA Hottried A., Schein J.E., Jones S.J.M., Marra M.A.,
RA Hottried A., Schein J.E., Jones S.J.M., Marra M.A.,
RA Hottried A., Schein J.E., Jones S.J.M., Marra M.A.,
RA Hottried A., Schein J.E., Jones J., Skalska U., Smailus D.E.,
RA Hottried A., Schein J.E., Jones J., Skalska U., Smailus D.E.,
RA Hottried A., Schein J.E., Jones J., Skalska U., Smailus D.E.,
RA Hottried A., Schein J.E., Jones J., John J., Skalska U., Smailus D.E.,
RA Hottried A., Schein J.E., Jones J., John J., John J., John J., John J., John J., John J., John J., John J.,
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Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
Schriml L.M.; Kanapin A., Matsuda H., Batalov S., Beisel K.W.,
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Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.B., Cousins S.,
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RA Blake J.A., Bradt D., Brusic V., Godzik A., Gough J.,
RA Gassterland T., Gartholdi M., Gissi C., Godzik A., Gough J.,
RA Gassterland T., Gustincich S., Hirokawa N., Jackson I.J., Jarvis B.D.,
RA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
RA Magashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,
Retrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,
RA Nagashima T., Schneider C., Semple C.A., Setou M., Shimada K.,
RA Rawasi T., Reed J.C., Redd D.J., Reid J., Ring B.Z., Ringwald M.,
RA Rawasi T., Redd J.C., Redd D.J., Reid J., Watanabe Y., Wells C.,
RA Wilming L.G., Wynshaw-Boris A., Yanagisawa M., Yang I., Yang L.,
RA Yana Z., Zavolan M., Zhu Y., Zimmer A., Carninci P., Hayatsu N.,
RA Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K.,
Hiroxani T., Waki K., Sasaki D., Shibata K., Shinagawa I.,
RA Yasunishi A., Yoshiho M., Waterston R., Lander E.S., Rogers J.,
RA Yasunishi A., Yashiho M., Waterston R., Lander E.S., Rogers J.,
RA Hara A., Hashizume W., Imotani K., Ishii Y., Itoh M., Kagawa I.,
RA Yasunishi A., Yashiho M., Waterston R., Lander E.S., Rogers J.,
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"Analysis of the mouse transcriptome 60,770 full-langth cDNAs.";
Nature 420:563-573(2002).
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                                                                                                                                                            clotting (By similari
SUBCELLULAR LOCATION:
ALTERNATIVE PRODUCTS:
                                                      Event=Alternative Name=HMW;
IsoId=008677-1; Sequence=Displayed;
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EMBL; D84415; BAA19742.1; -; mRNA.
EMBL; D805547; -; NOT ANNOTATED CDS; mRNA.
EMBL; BC01818; AAH18158.1; -; mRNA.
EMBL; BC01818; AAH18158.1; -; mRNA.
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the Euro
                                                                                                                                                                          SEQUENCE
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European
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PTM: Bradykinin is released from kininogen by plasma kallikrein.
SIMILARITY: Contains 3 cystatin-like domains.
                                          311
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                                                                                                                                82;
                                                                                                                                           Similarity
ALDM 374
                   PLGM 125
                                         VAGTKYVIEFIARETKCSKESNTELAEDCEIKHLGQSLDCNANVYMRPWENKVVPTVKCQ
                                                               VAGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQ
                                                                                     SGDDLVEALPKPCPGCPRDIPVDSPELKEVLGHSIAQLNAENDHPFYYKIDTVKKATSQV
                                                                                                        SGKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQV
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response; Plasma; Protease inhibitor; Repeat; Signal;
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Bioinformatics
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N-linked (GlcNAc. . .) (Potential).
N-linked (GlcNAc. . .) (Potential).
N-linked (GlcNAc. . .) (Potential).
Interchain (between heavy and light chains) (By similarity).
By similarity.
Gy Similarity.
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Cystatin-like 2.
Cystatin-like 3.
His-rich.
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RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausmer R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Pahey J., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalaka U., Smailus D.E.,
"Generation and initial analysis of more than 15,000 full-length human
"Generation and initial analysis of more than 15,000 full-length human
"Generation and initial analysis of more than 15,000 full-length human
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KNG1 RAT STANDARI
P08934; P08933;
01-NOV-1988 (Rel. 09, 0
01-NOV-1988 (Rel. 09, 1
13-SEP-2005 (Rel. 48, 1
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01-FEB-2005
01-FEB-2005
01-FEB-2005
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Submitted (DEC-2004) to the EMBL/GenBank/DDBJ databases.
Submitted (DEC-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; BC08815; AAH88155.1; -; mRNA.
GO; GO:0004869; F:cysteine protease inhibitor activity; IEA.
GO; GO:0004869; AAR; 47891 MW; 6D584348595166ED CRC64;
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
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                                                                                                                                                                                                                                                                                     375
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Last sequence up
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14; Mismatches
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J. Biol.
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MEDLINE-87137443; PubMed=3029068;
Kitagawa H., Kitamura N., Hayashida H., Miyata T., N
Kitagawa H., Kitamura N., Hayashida H., Miyata T., N
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Mammalia, Eutheria, Euarchontoglires, Glires, Rodentia, Sciurognathi,
Murcidea, Muridae, Murinae, Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kininogen-1 precursor (Contains:
Kininogen-1 light chain).
Name=Kng1; Synonyms=Kng;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kageyama R., Kitamura N., Ohkubo H., Nakanishi S.;
"Differing utilization of homologous transcription initiatirat K and T kininogen genes under inflammation condition."
J. Biol. Chem. 262:2345-2351(1987).
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"Structure and expression of the genes for major acute phase alpha protein (thiostatin) and kininogen in the rat.";
J. Biol. Chem. 262:9298-9308(1987).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=87250580; PubMed=2439509;
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
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                                                                                                                                                                                                                                                                                                                                                                                                                           SUBCELLULAR LOCATION:
ALTERNATIVE PRODUCTS:
Event=Alternative spl
                                                                                                                                                                 PTM: Bradykinin is released from kininogen by plasma kallikrein.
MISCELLANEOUS: Rats express four types of kininogens: the
classical HMW/LMW kininogens and two additional LMW-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FUNCTION: (1) Kininogens are inhibitors of thiol proteases;
                                                                                                 SIMILARITY: Contains 3 cystatin-like domains.
                                                                                                                                   classical HMW/LMW kininogens and kininogens: T-I and T-II.
                                                                                                                                                                                                                                                                   TISSUE SPECIFICITY:
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EMBL; M14369; AAA41485.1; A
EMBL; M14369; AAA41482.1; A
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PRODOM; PD001231; Prot inh 125A_B; 1.

PROSITE; PS00287; CYSTATIN; 2.

Alternative splicing; Blood coagulation;

Inflammatory response; Multigene family;

Repeat; Signal; Thiol protease inhibitor;
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RGD; 2980; kngl.

InterPro; IPR002395; Kininogen.

InterPro; IPR000010; Prot_inh_Cystat.

InterPro; IPR003243; Prot_inh_I25A_B.

PSTANCE_PRO0321; Cystatin; 3.
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Pfam; PF00031; Cystatin; 3.
SMART; SM00043; CY; 3.
PROSITE; PS00287; CYSTATIN; 1.
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"Structure and expression of two kin Biol. Chem. 385:295-301(2004).
EMBL; AY462057; AAR88631.1; -; mRNA MGI; MGI:3027157; Kng2.
GO; GO:0004869; F:cysteine protease
                                                                                                                                                   STRAIN=FUB/N;
MEDLINE=23306569; PubMed=15134344; DOI=10.1515/BC.2004.025;
Cardoso C.C., Garrett T., Cayla C., Meneton P., Pesquero J.
                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
Muridae; Murinae; M
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Q6S910 MOUSE PRELIMINARY;
Q6S910;
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Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                               NUCLEOTIDE
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InterPro; IPR0002395; Kininogen.
InterPro; IPR000010; Prot inh_cystat.
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MGI; MGI:3027157; Kng2.
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                                                                                                                                                                                                                                                                                                                                                                      PRINTS; PR00334; KININOGEN.
SMART; SM00043; CY; 3.
PROSITE; PS00287; CYSTATIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-23306569; PubMed-15134344; DOI=10.1515/BC.2004.025; Cardoso C.C., Garrett T., Cayla C., Meneton P., Pesquero J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bader M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
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  L 372
                                        ٦.
                                                                                                                      AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 122
                                                                                                                                                                          GDDLVEALPKPCPGCPRDIPVDSPELKEVLGHSIAQLNTENDHPFYFKIDTVKKATSQVV
                                                                                                                                                                                              GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                                                      AGTNYVI BFIARETKOSKESNTELTEDCEIKHLGQSLDCNANVYMRÞWENKVI ÞTVKCQA
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Pred. No. 2
                                                                                                                                                                                                                                                                                     Score 425;
Pred. No. :
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                                                                                          371
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InterPro; IPR000010; Prot\_inh\_cystat.
InterPro; IPR003243; Prot\_inh\_IZ5A\_B.
Pfam; PF00031; Cystatin; 3.
ProDom; PD001231; Prot\_inh\_IZ5A\_B; 1.
SMART; SM00043; CY; 3.
SMART; SM000287; CYSTMTIN; 2.
Acute phase; Bradykinin; Direct protein sequencing; Glycoprotein; Multigene family; Plasma; Protease inhibitor; Repeat; Signal; Thiol protease inhibitor; Vasoactive; Vasodilator.

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RESULT 13
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-KNT2 RAT
P08932;
01-NOV-1988
01-FEB-2005
13-SEP-2005
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13-SEP-2005 (Rel. 48, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
T-kininogen II precursor (Major acute phase protein) (Alpha-1-MAP)
(Thiostatin) [Contains: T-kininogen II heavy chain; T-kininogen II light chain].
kininogen II light chain].
Rattus norvegicus (Rat).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognat
Muroidea; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE-86008266; PubMed=2413019; Anderson K.P., Heath E.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=86008264; PubMed=2413018; Furuto-Kato S., Matsumoto A., Kitamura N., Nakanishi S.; Furuto-Kato S., Matsumoto A., Kitamura N., Nakanishi S.; "Primary structures of the mRNAs encoding the rat precursors for bradykinin and T-kinin. Structural relationship of kininogens with major acute phase protein and alpha 1-cysteine proteinase inhibitor."; J. Biol. Chem. 260:12054-12059(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       kininogens.";
J. Biol. Chem.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE
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PIR; B28055; B28055.
GlycoSuiteDB; P08932; -.
Ensembl; ENSRNOG00000030387; Rattus norvegicus.
                                                                                                                    EMBL; M11885; AAA41491.1;
EMBL; M11661; AAA41570.1;
                                                                                                                                                                                                                                                                                                   This Swiss-Prot entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict use as long as its content is in no way modified and this statuse as long as its content is in no way modified and this statuse as long as its content is in no way modified and this statuse as long as its content is in no way modified and this status as long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as its content is in no way modified and this status was long as long as its content is not way modified and this status was long as l
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SUBCELLULAR LOCATION: Extra:
TISSUE SPECIFICITY: Plasma.
INDUCTION: In response to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION: Kininogens are plasma glycoproteins with a number of functions: (1) as precursor of the active peptide bradykinin they effect smooth muscle contraction, induction of hypotension and increase of vascular permeability. (2) They play a role in blood coagulation by helping to position optimally prekallikrein and factor XI next to factor XII. (3) They are inhibitor of thiol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of the T-II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  kininogen I is raised.
PTM: As T-kinin is preceded by a Met instead of an !
is not released from its precursor by either tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SIMILARITY: Contains 3 cystatin-like domains. CAUTION: Ref.2 sequence differs from that shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in positions 180 and 181.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        classical HMW and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INDUCTION: In response to an inflammatory stimulant. II synthesis is induced and the plasma concentration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                same gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     260:12065-12071(1985).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ts express four types of kininogens:
LMW kininogens produced by alternat
and two additional LMW-like kininog
                                                                                                                    ; -; mRNA.
; ALT_FRAME;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Extracellular.
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Glires; Rodentia; Sciurognathi;
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MBL outstation -
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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler
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Mammalia, Eutheria, Euarchontoglires, Glires, Rodent
Muridae, Murinae, Rattus.
NCBI_TaxID=10116;
                                                                NUCLEOTIDE SEQUENCE.
TISSUE=Liver;
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Director MGC Project;
Submitted (DEC-2004) to the EMBL/GenBank/DDBJ (SMBL; BC088161; AAH88161.1; -; mRNA.
GO; GO:0004869; F:cysteine protease inhibitor InterPro; IPR000010; Prot inh cystat.
InterPro; IPR003243; Prot inh 125A_B.
The DEN0031; Cystatin; 3.
NUCLEOTIDE SEQUENCE.

PubMed=13679312; DOI=10.1095/biolreprod.103.020412;

Vonnahme K.A., Fernando S.C., Ross J.W., Ashworth M.D.,

Malayer J.R., Geisert R.D.;

Malayer J.R., Geisert R.D.;

Porcine endometrial expression of kininogen, factor XII

"Porcine endometrial expression of gilts.";

Biol. Reprod. 70:132-138(2004).

BMBL; AY321363; AAP85260.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                  Name=kng;
Sus scrofa (Pig)
Eukaryota; Metaz
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01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
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PROSITE; PS00287; CYSTATIN;
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Pfam; PF00031; Prot_inh_I25A_B; 1.
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